# Davis, Minh-Tam

To:

Schulwitz, Paul

Subject:

search for 09/700700

Thanks for your search result of SEQ ID NO:434 of 09/439313

However it seems that the search was not done against the parent cases of 09/439313, as requested, but rather against commecial data base, PGPUB and issued patents.

Could you please contact me ASAP because it is due this biweek.

Thanks.

MINH TAM DAVIS

305-2008

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# STIC Search Report Biotech-Chem Library

# STIC Database Tracking Number: 103561

TO: Minh-Tam Davis

Location: CM1/8A01&8E12

**Art Unit: 1642** 

Saturday, September 13, 2003

Case Serial Number: 09/700700

From: Paul Schulwitz

**Location: Biotech-Chem Library** 

CM1-6B06

Phone: 305-1954

paul.schulwitz@uspto.gov

# Search Notes

Examiner Davis,

See attached results.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services!

Paul Schulwitz
Technical Information Specialist
STIC Biotech/Chem Library
(703)305-1954



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### STIC-Biotech/Ch mLib

From:

Hutzell, Paula

Sent: To: Subject: Wednesday, September 10, 2003 1:57 PM STIC-Biotech/ChemLib; Davis, Minh-Tam FW Rush search request for 09/700700

## approved

----Original Message-

From:

Davis, Minh-Tam

Sent:

Wednesday, September 10, 2003 9:04 AM

To:

Hutzell, Paula

Subject:

FW: Rush search request for 09/700700

Could you approve the rush request since Christina is out of the office today? ThanksTam

----Original Message----

Fr m:

Davis, Minh-Tam

Sent:

Wednesday, September 10, 2003 9:02 AM Chan, Christina

To: Subject:

Rush search request for 09/700700

Please search SEQ ID NO:434 in the following parent cases of 09/439313 for priority date determination.

09439313 <snquery.pl?APPL ID=09439313>

is a continuation in part of <u>09352616</u> < snquery.pl?APPL ID=09352616>

Which is a continuation in part of 09288946 < snquery.pl?APPL ID=09288946>

Which is a continuation in part of 09232149 < snquery.pl?APPL ID=09232149>

Which is a continuation in part of <u>09159812 < snquery.pl?APPL\_ID=09159812></u>

Which is a continuation in part of 09115453 < snquery.pl?APPL ID=09115453>

Which is a continuation in part of 09030607 < snquery.pl?APPL ID=09030607>

Which is a continuation in part of 09020956 < snquery.pl?APPL ID=09020956>

Which is a continuation in part of <u>08904804</u> <snguery.pl?APPL\_ID=08904804>

Which is a continuation in part of <u>08806099</u> <snguery.pl?APPL\_ID=08806099>

Thank you.

MINH TAM DAVIS

<u>ART UNIT 1642, ROOM 8A01, MB 8E12</u>

305-2008

Searcner:	
Phone:	
Location:	
Date Picked Up:	9/12
Date Completed:	9/13
Searcher Prep/Rev	iew:
Clerical:	
Online time:	

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
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VENDOR/COST (where ap	plic.)
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DIALOG:	
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DRLink:	
Lexis/Nexis:	
Sequence Sys.:	
WWW/Internet:	
Other (specify):	

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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

DB Length

Query

Score

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RESULT 1 AR261002 LOCUS O 0 0 0000000000 (without alignments) 9145.616 Million cell updates/sec September 13, 2003, 01:49:31; Search time 2165 Seconds 1 ttttaaaaataagcatttagt......ataaagtacccatgtcttta 5777422 GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd. 2888711 segs, 20454813386 residues Total number of hits satisfying chosen parameters: Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries - nucleic search, using sw model IDENTITY\_NUC Gapop 10.0 , Gapext 1.0 em\_htgo\_other:\* em\_htg\_other:\* US-09-352-616A-434 484 em\_htgo\_hum:\* em\_htgo\_mus:\* seq length: 0 seq length: 2000000000 em\_vi:\* em\_htg\_hum:\* em\_htg\_mus:\* em\_htg\_pln:\* em\_htg\_inv:\* em\_htg\_rod:\* em\_htg\_mam:\* em\_htg\_vrt:\* gb\_ro:\*
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Sequence Sequence

AR244194 SAR237207 AR237207 AR2760700 AR2706334 AX10625 AX207141 BD070260 AX337002

AR260700 AR278231 AX106334 AX140625 AX200485 AX267141 BD070260

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AR237233 AR260726 AR278257 AX106360 AX140651 AX200511

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AX140944 AX200804 AX267460 AR244306 AR237420 AR260913

Sequence

AR261002

Description

Unknown.
Unknown.
Unclassified.
1 (bases 1 to 484)
Mashiki.2. and Harada.J.
Negative pressure control apparatus for engine mounted in vehicle Patent: US 6321716-A 434 27-NOV-2001;
Location/Qualifiers Unknown. DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL FEATURES

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PAT 29-JAN-2003

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ALIGNMENTS

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AR261002 AR261002.1 GI:28071765

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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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100.0%; Pred. No. 4.5e-105;
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Sequence 434 from Patent WO0151633.
AX200804
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QY         241 AGCTAGTCTATCAGCATCTGACAGTGATTGGATGGTTCTCAGAACCATTCACCCAGA 300           Db         241 AGCTAGTCTATCACAGCTCGACAGGTGATTGTTCTCAGAACCATTTCACCCAGA 300           CA         301 CAGCTGTTCTATCATCATTAGTTTGGTTCTCTACATGCATAACAACCC 360           Bb         301 CAGCCTGTTTCTATCTTTAATAATTAGTTTGGTTCTTACATGCATAACAACCC 360           CAGCCTGTTTCTATCCTGTTTAATAATTAGTTTGGTTCTTCTACATGCATACAAACCC 360           CAGCCTGTTTCTAATCATTAATAAATTAGTTTGGGTTCTTCTACATGCATACAAACCC 360           CAGCCTGTTTCTAATCATTAATAAATTAGTTTGGGTTCTTCTACATGCATACAAACCC 360           CA         361 TGCTCCAATCTGTCACATAAAAATTAGTTTGAGTTTTGAACACCCCCCCC	RESULT 6 AX267460 AX2	

ACCESSION AR260913 VERSION KEYWORDS SOURCE Unknown. ORGANISM Unclassified.  NGANISM Unclassified.  AUTHORS AUTHORS AUTHORS AUTHORS AUTHORS Location/Qualifiers Location/Qualifiers Source Authors Auth	Query Match         97.5%;         Score 472;         DB 6;         Length 2984;           Best Local Similarity         99.6%;         Pred;         No. 2.7e-102;           Matches 484;         Conservative         0;         Mismatches         0;         Indels         2;         Gaps         1;           Qy         1         TTTTAAAATAAGCATTTAGTGCTCAGTCCTACTGAGTACTCTTCTCCCCTCCTCG         60         111111111111111111111111111111111111	TGTTGCAAAAAAAAAAAAAGTCTTTGTTTAAATTACTTGGTTTGTGAATCCATCTT	Db   2653 GCTTTTCCCCATTGGAACTAGTCATTAACCCATCTCGAACTGGTAGAAAACATCTGA 2712	OY 299 GACAGCCTGTTTCTATCCTGTTTAATAATTAGGTTCTCTACATGCATAACAAAC 358	Qy 359 CCTGCTCCAATCTGTCACATAAAAGTCTGTGAACTTTAAGTCAGCACCCCCACCAA 418 	QY 419 ACTITATITITCTATGTGTTTTTTGCAACATATGAGTGTTTTGAAAATAAGTACCCATG 478 	Oy 479 TCTTA 484 	T 10	PAI	Σ	REFERENCE 1 (bases 1 to 2984) AUTHORS Xu,J., Dillon,D.C., Mitcham,J.L., Harlocker,S.L., Jiang,Y., Kalos,M.D., Fanger,G.R., Retter,M.W., Stolk,J.A., Day,C.H., Vedvick,T.S., Carter,D., Li,S.X., Wang,A., Skeiky,Y.A.W., Hepler,W.T. and Henderson,R.A. TITLE Compositions and methods for the therapy and diagnosis of prostate
37 43	ORGANISM Unknown.  REFERENCE 1 (bases 1 to 2984)  AUTHORS Xu,J., Dillon,D.C. and Mitcham,J.L.  TITLE Compounds for immunotherapy of prostate cancer and methods for tent: use their use  JOURNAL Patent: US 6465611-A 335 15-OCT-2002;  FEATURES Location/Qualifiers  Source /Organism="unknown"  BASE COUNT 837 a 659 c 645 g 842 t 1 others	<pre>Duery Match</pre>	DD 24/3 TTTTAAATAAGCATTTAGTGCTCCACTGCTACTGAGTACTCTTCTCTCCCTCTG 2532  Qy 61 AATTTAATTCTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTG 120  DD 2533 AATTTAATTCTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATG 2592	OY 121 TGTTGCAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGATTGTGAATCCATCTT 178 	OY 179 GCTTTTCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAACATCTGA 238	OY 239 AGAGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTCACCA 298	Oy 299 GACAGCCTGTTTCTATCCTGTTTAATAAATTAGTTTGGGTTCTCTACATGCATAACAAC 358	OY 359 CCTGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCCACCAA 418 	Qy 419 ACTITATITICTATGTTTTTTGCAACATATGAGTGTTTTGAAAATAAAGTACCCATG 478 	Oy 479 TCTTA 484  L       Db 2953 TCTTA 2958	RESULT 9 ARAS60913 LOCUS AR260913 2984 bp DNA linear PAT 29-JAN-2003 DEFINITION Sequence 335 from patent US 6321716.

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61 AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATTG 120
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Manmalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 2.7e-102;
0; Mismatches 0;
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99.6%; Pred. No. 2.7e-102;
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/Organism="Homo sapiens"
/Mol_type="genomic DNA"
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99.6%;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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Pred. No. 2.7e-102;
0; Mismatches 0;
         US 6512094-A 335 28-JAN-2003;
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/organism="unknown"
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//organism="Homo sapiens"
/mol_type="genomic DNa"
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                                                    CCTGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCCACCAA
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           GCTTTTTCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAACATCTGA
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
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Kalos,M.D., Fanger,G.R., Retter,M.W., Stolk,J.A., Day,C.H.,
Vedvick,T.S., Carter,D., Li,S.X., Wang,A., Skeiky,Y.A., Hepler,W.T.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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23-SEP-1998;
15-JAN-1999;
15-JAN-1999;
09-APR-1999;
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                                                                                                     September 12, 2003, 23:52:20; Search time 225 Seconds (without alignments) 5806.791 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N. Geneseq_19Jun03:*

| SIDS1/gcgdata/geneseq/geneseqn-embl/NA1980.DAT:*
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               GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd
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Maximum Match 100%
Listing first 45 summaries
                                                                           nucleic search, using sw model
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The present invention describes isolated polypeptides, comprising an and polymucleotides encoding them have cytostatic activity and can be used in vaccines and in gene therapy. The polypeptides and in open therapy. The polypeptides and polymucleotides encoding them, antigen presenting cells which express comprising them can be used for inhibiting the development of prostate cancer in a patient. The polypeptides and vaccines cancer in a patient. The polypeptides can be used to generate antibodies the polymucleotides encoding the polypeptides can be used to generate antibodies the polymucleotides encoding the polypeptides can be used as a probe or computation of modulate the expression of the polypeptides. AAA06241 to AAA06691 and the present invention.
                                                                                                                                                                                                                                                                                                                                                 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                      181 TTTTCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAACATCTGAAG 240
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                                                                 cancer
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0
                                                             prostate or protein -
                                                                                                                                                                                                                                                                                                              Score 484; DB 21; Length 484;
Pred. No. 1.4e-116;
Mismatches 0; Indels 0;
           Mitcham JL;
                                                                                                                                                                                                                                                                                Sequence 484 BP; 137 A; 101 C; 73 G; 173 T; 0 other;
                                               New polypeptide useful for treating and diagnosing comprises an immunogenic portion of prostate tumor
          ٦,
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       Yuqin J,
                                                                                  Claim 1; Page 250; 263pp; English.
                                                                                                                                                                                                                                                                                                                                      ;
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     SL,
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   Harlocker
                        WPI; 2000-171268/15
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 Dillon DC,
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AAS63862 standard; cDNA; 484 BP.

(first entry)

29-JAN-2002

AAS63862;

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The invention relates to isolated prostate-specific polypeptides, polypeptides, fusion proteins of the polypeptides, antibodies raised against the polypeptides (or antigenic epitopes derived from them) and antigen researching cells expressing the polypeptides. The antibodies are useful for detecting the presence of the antigen-presenting cells are useful for stimulating and/or expanding of cancer especially prostate cancer. The polypeptides, polynucleotides and T cells specific for a tumour protein, and for inhibiting the development polynucleotide and/or polypeptide are useful for stimulating the development polynucleotide and/or polypeptide are useful for stimulating an immune response, and for treating cancer. The oligonucleotide is useful for polynucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATTG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New human prostate-specific polypeptides and polynucleotides useful for the diagnosis and treatment of cancer, especially prostate cancer -
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Carter D;
                                       cancer; ss; cytostatic; immunostimulant; tumour.
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                                                                                                                                                                                                                                                                                                                               Dillon DC, Mitcham JL, Harlocker SL, Jiang Y, GR, 'Retter MW, Stolk JA, Day CH, Vedvick TS, Wang A, Skeiky YAW, Hepler WT, Henderson RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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Pred. No. 1.4e-116;
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          Human prostate cDNA sequence #398
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                                                                                                                                                                              ; 2000us-0568100
; 2000us-057037
; 2000us-053793;
; 2000us-0636215;
; 2000us-0651236;
; 2000us-0651279;
; 2000us-0651279;
; 2000us-0651279;
                                                                                                                                     27-MAR-2001; 2001WO-US09919.
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100.0%;
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Best Local Similarity
                                       prostate
                                                                                      WO200173032-A2.
                                                                                                                                                                                                    13-JUN-2000;
27-JUN-2000;
10-AUG-2000;
29-AUG-2000;
06-SEP-2000;
02-OCT-2000;
10-OCT-2000;
                                                              Homo sapiens.
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12-MAY-2000;
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                                      Human;
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Sequence 484 BP; 137 A; 101 C; 73 G; 173 T; 0 other;
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18-NOV-1999;
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360
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                                                                                                           361 IGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCCACCAAAC 420
                                                                                                                                            CAGCCTGTTTCTATCCTGTTTAATAAATTAGTTTGGGTTCTCTACATGCATAACAAACCC
                                                        TGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCCACCAAC
                                                                                                                                 TITATITITCTATGTGTTTTTTGCAACATATGAGTGTTTTGAAAATAAAGTACCCATGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New polynucleotide encoding a prostate-specific protein, for diagnosing, monitoring and treating prostate cancer in a patient and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Harlocker SL, Jiang Y, Reed SG;
Retter MW, Stolk JA, Skeiky YAW;
                                                                                                                                                                                                                                                                                                                                             prostate cancer; prostate-specific; diagnosis; vaccine;
                                                                                                                                                                                                                                                                                                                       Human prostate-specific cDNA sequence 22595
                                                                                                                                                                                                                                                                                                                                                         therapy; metastasis; ss.
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                                                                                                                                                                                                                                                     AAH93769 standard; cDNA; 484
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JAN-2000; 2000US-0483672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Xu J, Dillon DC, Mitcham
Kalos MD, Fanger GR, Day
Wang A, Meagher MJ;
                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for use in vaccines -
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TTTA 484
                                                                                                                                                                                                                                                                                                                                                         cytostatic; gene
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                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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                                                                                                                                                          AGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCAGA
                                                                                                                                                                                                                                                                                                                                                                                                                   CAGCCTGTTTCTATCCTGTTTAATAAATTAGTTTGGGTTCTCTACATGCATAACAAACCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITATITITICIATGITITITICCAACATATGAGTGTTTTGAAAATAAAGTACCCATGTC
                                                                    1 TITIAAAATAAGCATITAGIGCICAGICCCIACIGAGIACICTITCICCCCCCCCCTCIG
                                   Gaps
                                                                                                                                                                                                                                                                                                                                                               Reed SG;
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                                  ö
Length 484;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; prostate cancer; therapy; diagnosis; cat eye syndrome; chromosome 22q11.2; prostate-specific protein; chromosome 1; prostate specific antigen; PSA; ss.
                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Harlocker SL, Jiang Y,
Day CH, Skeiky YAW,
Score 484; DB 22;
Pred. No. 1.4e-116;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human prostate-specific cDNA sequence 22595.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Xu J, Dillon DC, Mitcham JL, H. Kalos MD, Retter MW, Stolk JA,
100.0%; Sc
100.0%; Pr
tive 0;
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99US-0443686
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                                  Matches 484; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CORI-) CORIXA CORP.
   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TTTA 484
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us-09-352-616a-434.rng

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The present invention describes an isolated polypeptide (I) comprising at least an immunogenic portion of a prostate tumour antigion protein or trs variant. (I) have cytostatic activity and can be used in vaccine production. (I), prostate tumour antigon polynocleotides, an antigen protein or presenting cell (APC e.g. a dendrile cell) that expresses (I), and a cevelopment of cancer in a patient. Antibodies specific for inhibiting the specific proteins and oligonucleotides that hybridise to a colynucleotide that encodes a prostate specific protein are useful conferesion the progression of a cancer. Conferession the progression of a cancer or anonitoring the progression of a cancer, especially prostate cancer.

AH02422 to AH12872, AAB74798 to ABB74821 and AAB74830 are sequences were conferenced.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGTTGCAAAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTGC 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         241 AGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCAGA 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                umour antigen; prostate tumour; therapy; diagnosis;
immunogenic; cytostatic; vaccine; ss.
                                                                                                                                                                                                                                                         Prostate specific protein and its encoding polynucleotide, useful for
the treatment and diagnosis of prostate cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 484; DB 22;
1100.0%; Pred. No. 1.4e-116;
ive 0; Mismatches 0;
                                                                                                                                                                                                                   Cheever MA;
                                                                                                                                                                                                                                                                                          Claim 4; Page 256; 276pp; English.
                                       tumour antigen;
                                                                                                                                                                                                                 SG,
                                                                                                                                                                  99US-0157455.
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                                                                                                                                                                                    (CORI-) CORIXA CORP
                                                                                                                                                                                                                                  WPI; 2001-245062/25.
                                                                                                                                                                                                             Skeiky YAW,
                                    Human; prostate t
prostate cancer;
                                                                                            WO200125272-A2
                                                                                                                                        04-OCT-2000;
                                                                                                                                                              04-OCT-1999;
                                                                                                                  12-APR-2001
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                                                                                         The present invention describes an isolated polypeptide (PI) comprising variant. Also described are polynucleotides (NI) encoding (PI). Or its (NI) have cytostatic activity and can be used in vaccine production. The polypeptides, nucleic activity and can be used in vaccine production. Prostate specific activity Pr12P, PT1AP, PT75P and Present in a genomic region on chromosome 22q11.2 known as the Cat Eye Syndrome region. Prostate specific antigen (PSA) P501S was located on chromosome 22q11.2 known as the Cat Eye Syndrome chromosome I. AAH84671 to AAH8543 and AAG99000 to AAG99077 represent of the present invention.
                                                                                                                                                                                                                                                                                                                                                                            AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTTTCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAAACATCTGAAG 240
                                                                                                                                                                                                                                                                                                                 1 TITTAAAATAAGCAITTAGTGCTCAGTCCCTACTGAGTACTCTTTCTCTCCCCCTCTG 60
                                                                                                                                                                                                                                                                                                                               300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CAGCCTGTTTCTATCGTGTTTAATAAATTAGTTTGGGTTCTCTACATGCATAACAAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isolated polypeptide comprising at least an immunogenic portion of prostate-specific protein, useful in the diagnosis and therapy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCACCAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITATITITICIATGIGITITITIGCAACAIAIGAGIGITITIGAAAATAAAGIACCCAIGIC
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                                                                                                                                                                                                                                                                                Length 484;
                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                             Sequence 484 BP; 137 A; 101 C; 73 G; 173 T; 0 other;
                                                                                                                                                                                                                                                                       100.0%; Score 484; DB 22;
100.0%; Pred. No. 1.4e-116;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prostate tumour antigen cDNA sequence for 22595.
                                                                  Claim 5; Page 273; 325pp; English.
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                                                                                                                                                                                                                                                                            Best Local Similarity 100.
Matches 484; Conservative
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                                             prostate cancer
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|TTA 484
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Length 484; Indels S

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The present invention provides prostate-specific coding sequences and their encoded proteins. These can be used in the diagnosis and treatment of cancers, particularly prostate cancer. The present sequence is a CDNA described in the invention.
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                                                                                                                                                                                                                                                         61 AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATTG
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                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prostate cancer; vaccine; gene therapy; cytostatic; fusion protein; immunogen; cancer; prostate specific antigen; PSA; prostatic acid phosphatase; PAP; prostate specific membrane antigen; PSMA; gene; ss.
                                   New prostate-specific polynucleotides for diagnosing and treating diseases, in particular prostate cancer, and as markers for the progression of cancer
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                                                                                                                                                                    Length 484;
                                                                                                                                                                                      Indels
 Henderson RA;
                                                                                                                                               Sequence 484 BP; 137 A; 101 C; 73 G; 173 T; 0 other;
                                                                                                                                                                  Score 484; DB 24;
Pred. No. 1.4e-116;
                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prostate cancer therapy associated cDNA #413.
                                                                                                                                                                                     0; Mismatches
 Hepler WT,
                                                                            Claim 1; SEQ ID NO 434; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                   100.0%;
100.0%;
 Skeiky YAW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACA59670 standard; cDNA; 484
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                                                                                                                                                                                      Matches 484; Conservative
                  WPI; 2002-255649/30.
                                                                                                                                                                             Similarity
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 SX, Wang A,
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 TTTATTTTTCTATGTGTTTTTTGCAACATATGAGTGTTTTGAAAATAAAGTACCCATGTC
          cancer; prostate cancer; vaccine; cytostatic; immunostimulant;
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Carter D;
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                                                                                                                                                                   Human 22595 cDNA sequence SEQ ID NO 434.
                                                                                                         BP.
                                                                                                                                                                                                                                                                                             9705-0806099.
9705-0904804.
9805-0030607.
9805-0115453.
9805-0115453.
9905-0159812.
9905-0282149.
9905-043913.
9905-043913.
9905-0443913.
200005-0508100.
200005-050805.
                                                                                                        ABL95233 standard; cDNA; 484
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                                                                                                                                                (first entry)
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HARLOCKER S L.
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DAY C H.
VEDVICK T S.
                                                                                                                                                                                    Human; cancer; prosu
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DILLON D C.
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KALOS M D.
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|TTTA 484
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WANG A.
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09-MAY-2000;
12-MAY-2000;
13-JUN-2000;
27-JUN-2000;
27-AUG-2000;
29-AUG-2000;
                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                            12-JAN-2001;
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13-JUL-1999;
12-NOV-1999;
18-NOV-1999;
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09-FEB-1998
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                  421
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421
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(MITC/)
(HARL/)
(JIAN/)
(FALC/)
(FANG/)
(RETL/)
(STOL/)
(DAYC/)
(VEDV/)
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(WANG/)
(SKEI/)
(HEPL/)
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                                                                                               ABL95233
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180

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241 AGCTAGICIATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCAGA 300
                                                                                                                                                                                                                                                                                                                            TGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCCACCAAAC 420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; prostate cancer; prostate cancer antigen; detection; diagnosis; neuroprotective; cytostatic; cardioactive; immunomodulatory; muscular; vulnerary; gastrointestinal; nephrotropic; antiinfective; gynaecological; antibacterial; gene therapy; neural; immune; reproductive; renal; gastrointestinal; pulmonary; cardiovascular; proliferative disorder; wound; infectious disease; ss.
                                                                             TTTTTCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAAAACATCTGAAG
                                                                                                                                                                                                                                                                       Prostate cancer associated gene sequences, referred to as prostate cancer antigens, useful for treatment, prevention, and diagnosis of disorders such as prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human prostate cancer antigen nucleotide sequence SEQ ID NO:258.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HUMA-) HUMAN GENOME SCI INC (ROSE/) ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-MAR-2000; 2000WO-US05988.
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481 TTTA 484
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes a fusion protein comprising at least one amino defined in the specification, or sequences having at least 70 or 90 sequences in the specification, or sequences having at least 70 or 90 sequences defined in the specification. The 35 sequences defined in the USPTO of 64 fined in the specification. The fusion protein, composition and composition and particularly prostate cancer. The proteins are useful as markers to indicate the presence or absence of cancer. This sequence of cancer therapy associated obna.

Once: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly cancer or cancer. The sequence of the sequence of the presence of the proteins are useful as markers to composite the presence of the proteins are useful as markers to composite the presence of the patent of the patent of the patent of the patent of the presence of the patent                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New fusion protein comprising prostate-specific polypeptides, or its immunogenic portions, useful for diagnosing, preventing and/or treating cancer, particularly prostate cancer
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Hural J;
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100.0%; Score 484; DB 25; Length 484;
Best Local Similarity 100.0%; Pred. No. 1.4e-116;
Matches 484; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dillon DC, Mitcham JL, Harlocker SL, Jiang Y, GR, Retter MW, Stolk JA, Day CH, Vedvick TS, Wang A, Skeiky YAW, Hepler WT, Henderson RA, I PD, Houghton RL, Y De Bassols CV, Foy TM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 484 BP; 137 A; 101 C; 73 G; 173 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      segdata.uspto.gov/sequence.html?DocID-US20020192763.
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                                                                               29-JUN-2001; 2001US-0895793.
                                                                                                                       17-APR-2000; 2000US-157455P.
04-OCT-2000; 2000US-0679272.
28-MAR-2001; 2001US-0822827.
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MCNEILL P D.
HOUGHTON R L.
Y DE BASSOLS C V.
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HEPLER W T.
HENDERSON R A.
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MITCHAM J L.
HARLOCKER S L
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KALOS M D.
KALOS M D.
RETTER M W.
STOLK J A.
DAY C H.
VEDVICK T S.
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  US2002192763-A1.
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WANG A.
                                          19-DEC-2002.
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Mcneill PD,
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Fanger GR,

Xu J,

(HOUG/) (DBAS/) (FOYT/) (HURA/) (MCNE/)

(HARL/)
(JIAN/)
(KALO/)
(FANG/)
(RETT/)

(STOL/) (DAYC/) (VEDV/)

(LISX/) (WANG/) (SKEI/) (HEPL/)

(DILL/) (MITC/)

(XCCOX)

Claim 1; Page 781; 2338pp; English

δ QQ

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Homo sapiens
                       WO9964594-A2
                                                                    .0-JUN-1999;
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                                             16-DEC-1999.
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     AAP15566 to AAF16505 encode the human prostate cancer associated proteins, called prostate cancer antigens, given in AAB56363 to AAB57302. The prostate cancer antigens can have neuroprotective, cytostatic, cardioactive, immunomodulatory, muscular, vulnerary, gastrointestinal, nephrotropic, antiinfective, gynaecological and antibacterial activities, and can be used in gene therapy. The prostate cancer antigen polynucleotides may be used for detection of prostate cancer, chromosome identification, as chromosome markers, and for numerous other diagnostic or research purposes. The prostate cancer antigens may be used to treat disorders such as neural, immune, muscular, reproductive, gastrointestinal, pulmonary, cardiovascular, renal, and proliferative disorders, wounds, and infectious diseases. AAF16506 to AAF16514 to
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                                                                                                                                                                                                                                1;
                                                                                                                                                                                                        21; Length 755;
                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                Sequence 755 BP; 217 A; 161 C; 128 G; 246 T; 3 other;
                                                                                                                                                                                                      Score 473; DB 21;
Pred. No. 1.2e-113;
0; Mismatches 0;
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ID AA297473 standard; cDNA; 2051
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722 CTTTA 726
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a prostate cancer specific nucleic acid sequence. The invention relates to a method for diagnosing cancer, tumour progression, hyperproliferative cell growth or accompanying biological and physical manifestations. The method involves contacting the biological sample with a probe that comprises a sequence capable of hybridising to any of the 339 nucleotide sequences given in the products and methods of the invention can be used for the diagnosis, prognosis, and treatment of cancer, tumour progression, hyperproliferative cell growth, and accompanying physical and biological manifestations. They can be used particularly for prostatic disorders such as metastatic prostate cancer, localised prostate cancer, or benign
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9
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                                                                                                                                                                                                                                                                            New isolated prostate cancer specific nucleic acids, used to products for the diagnosis and treatment of cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                             Monahan JE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 472; DB 21;
Pred. No. 2.9e-113;
0; Mismatches 0;
                                                                                                                                                             DW,
                                                                                                                                                          Ford
                                                                                                                                                                                                                                                                                                                                               Claim 2; Page 209-210; 212pp; English.
                                                                                                                                                               Endege WO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97.5%;
99.6%;
                                         98US-0088877
99US-0088877
99WO-US13181
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Steinmann KE, Zhang J;
                                                                                                                 (CHIR ) CHIRON CORP.
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2712

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2533 AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATG 2592
                                                                                                                                                                                     2593 TGTTGCAAAAAAAAAAAAGTGTCTTGTTTAAAATTACTTGGTTTGTGAATCCTTT
                                                                                GCTTTTTCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAAACTCTGA
                                                                                              AGAGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCA
                                                                                                                                          TGTTGC---AAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTT
                                                                                                                                                                          299 GACAGCCTGTTTCTATCTTTAATAATTAGTTTGGGTTCTCTACATGCATAACAAAC
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GR, Retter MW, Stolk JA, Day CH, Vedvick TS,
Wang A, Skeiky YAW, Hepler WT, Henderson RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human prostate cDNA sequence #325.
                                                                                                                                                                                                                                                                                                                                                                                         AAS63773 standard; cDNA; 2984 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; 2000us-0536B57.
2000us-056B100.
2000us-057037.
2000us-06057B3.
2000us-06057B3.
2000us-061215.
2000us-0651236.
2000us-0657279.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001WO-US09919
                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                   2953 TCTTTA 2958
                                                                                                                                                                                                                                                                                                               TCTTTA 484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-639232/73.
P-PSDB; AAU69771.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GR, Kelle,
Wang A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200173032-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-MAY-2000;
13-JUN-2000;
27-JUN-2000;
10-AUG-2000;
29-AUG-2000;
06-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-MAR-2000;
09-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-OCT-2000;
                                       121
                                                                                  179
                                                                                                                                                                                                                                                                                                               479
                                                                                                                                                                                                                                                                                                                                                                                                                AAS63773;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fanger
Li SX,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Xu J,
                                                                                                                                                                                                                                                                                                                                                                     RESULT 11
                                                                                                                                                                                                                                                                                                                                                                             AAS63773
                                                       Db
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                                                                                                                                                                                                                                                                                                                                 Db
ACTITALITITCIANGTGTTTTTTGCAACATATGAGTGTTTTGAAAATAAAGTACCCATG 375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes isolated polypeptides, comprising an and polymelectides encoding them have cytostatic activity and can be used in vaccines and in gene therapy. The polypeptides and be polymelectides and in gene therapy. The polypeptides and be polymelectides encoding them, antigen presenting cells which express comprising them can be used for inhibiting the development of prostate ancer in a patient. The polypeptides can be used to generate antibodies the polymelectides encoding the polypeptides can be used to generate antibodies the polymelectides encoding the polypeptides can be used as a probe or anti-idiotypic antibodies for passive immuno therapy. A portion of the polymelectides encoding the polypeptides can be used as a probe or AAV80000 to AAV8020 represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTG 120
                                                                                                                                                                    Human immunogenic prostate tumour protein cDNA sequence SEQ ID NO:335.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 TITIAAAATAAGCATTIAGTGCTCAGTCCCTACTGAGTACTCTTTCTCTCCCCTCCTG
                                                                                                                                                                                            Human; prostate cancer; diagnosis; tumour; gene therapy; detection; immunogenic; cytostatic; vaccine; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            \operatorname{polypeptide} useful for treating and diagnosing prostate cancer prises an immunogenic portion of prostate tumor protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 2984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          Mitcham JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2984 BP; 837 A; 659 C; 645 G; 842 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 472; DB 21;
Pred. No. 3.2e-113;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Xu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 50; Page 207-208; 263pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Yuqiu J,
                                                                                                AAA06564 standard; cDNA; 2984 BP.
                                                                                                                                                                                                                                                                                                                     98US-0115453.
98US-0116134.
98US-0159812.
99US-0232149.
99US-0232880.
                                                                                                                                                                                                                                                                                                 99WO-US15838
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99.6%;
                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                Dillon DC, Harlocker SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 99.6
Matches 484; Conservative
                                               369
                         TCTTTA 484
                                                                                                                                                                                                                                                                                                                                                                                                            (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-171268/15.
                                      TCTTTA
                                                                                                                                                                                                                                                 WO200004149-A2
                                                                                                                                                                                                                              Homo sapiens,
                                                                                                                                              13-JUN-2000
                                                                                                                                                                                                                                                                                               14-JUL-1999;
                                                                                                                                                                                                                                                                        27-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                            15-JAN-1999;
09-APR-1999;
                                                                                                                                                                                                                                                                                                                     14-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                 5-JAN-1999
                                                                                                                                                                                                                                                                                                                                          23-SEP-1998
                                                                                                                                                                                                                                                                                                                                                        23-SEP-1998
                      479
                                           374
  434
                                                                                                                         AAA06564;
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                                                                              RESULT 10
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human prostate-specific polypeptides and polynucleotides useful for diagnosis and treatment of cancer, especially prostate cancer

Kalos MD; Carter D;

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WO200151633-A2
        Homo sapiens
                                                                                                                                                         19-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2653 (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wang A,
    ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2773 GACAGCCTGTTTCTATCCTGTTTAATAATTAGTTTGGGTTCTCTACATGCATAACAAC 2832
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2833 CCTGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCCCCACCAA 2892
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to isolated prostate-specific polypeptides, fusion proteins of the polypeptides, antibodies raised against the polypeptides (or antigonic epitopes derived from them) and antigon-presenting cells expressing the polypeptides. The antibodies are useful for detecting the presence of cancer, especially prostate cancer. The polypeptides, polynucleotides and the antigen-presenting cells are useful for stimulating and/or expanding T cells specific for a tumour protein, and for inhibiting the development of cancer especially prostate cancer. Compositions comprising the polynucleotide and/or polypeptide are useful for stimulating an immune response, and for treating cancer. The oligonucleotide is useful for detecting cancer. The present sequence is a prostate specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2473 TITIAAAATAAGCATTTAGTGCTCAGTCCCTACTGAGTACTCTTTCTCTCCCCTCCTC 2532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2592
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  239 AGAGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCA 298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GACAGCCTGTTTCTATCCTGTTTAATAAATTAGTTTTGGGTTCTCTACATGCATAACAAAC 358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCTGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCCACCAA 418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GCTTTTTCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAACATCTGA 238
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     419 ACTITATITITICIATGTGTTTTTTGCAACATATGAGGTGTTTTGAAAATAAAGTACCCATG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 2984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2984 BP; 837 A; 659 C; 645 G; 842 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
0
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Pred. No. 3.2e-113;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human prostate-specific cDNA sequence P705P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytostatic; gene therapy; metastasis; ss.
                                    Page 335-336; 579pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 99.6 Matches 484; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       111111
2953 TCTTTA 2958
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                                    Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH93680;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         encode
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9
                                                                                                                                                                                                                                                                                                                                                                                                                                            and can be used in vaccine proteins (II). (I) and (II) have cytostatic activity, and can be used in vaccine production and gene therapy. (I), (II), antibodies to (II), fusion proteins comprising (II), and isolated T cells prepared using (I) or (II) are used treat cancer in a patient. The cancer that is diagnosed or treated is particularly prostate cancer. (I) and (II) can be used for monitoring the progression of cancer in a patient. (I) and (II) can be used for monitoring the progression of cancer in a method for monitoring the progression of cancer in a patient. (I) and (II) can be used to improve diagnostic and thermore.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (I) and (II) can also be used to improve diagnostic and therapeutic methods for prostate cancer. They can indicate the level of metastasis as well as the prostate volume. AAH93357 to AAH93944 and AAM01115 to AAM01318 represent polynucleotide and amino acid sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2473 TITITAAAATAAGCATITAGTGCTCAGTCCCTACTGAGTACTCTTCTCTCCCCTCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2533 AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGTTTATATTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                121 TGTTGC--AAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCTTTTTCCCCCATTGGAACTAGTCATTAACCCCATCTCTGAACTGGTAGAAAAACATCTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGAGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GACAGCCTGTTTCTATCCTGTTTAATAATTAGTTTGGGTTCTCTACATGCATAACAAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2773 GACAGCCTGTTTCTATCCTGTTTAATAATTAGTTTGGGTTCTCTACATGCATAACAAAC
                                                                                                                                                                                                                                                                                                 New polynucleotide encoding a prostate-specific protein, for diagnosing, monitoring and treating prostate cancer in a patient and for use in vaccines -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                   SG;
YAW;
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                                                                                                                                                     SL, Jiang Y, Reed
Stolk JA, Skeiky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 2984;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 472; DB 22;
Pred. No. 3.2e-113;
0; Mismatches 0;
                                                                                                                                                     Harlocker S
Retter MW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 333-334; 543pp; English.
                                                                                                                                                   Mitcham JL,
3R, Day CH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97.5%;
99.6%;
16-JAN-2001; 2001WO-US01574
                                                 14-JAN-2000; 2000US-0483672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 99.6
Matches 484; Conservative
                                                                                                                                                                             Fanger GR,
                                                                                                                                                                        3 MD, Fauyer
A, Meagher MJ;
                                                                                                 (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                  WPI; 2001-425873/45
                                                                                                                                                   Dillon DC,
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298

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299 GACAGCCTGTTTCTATCCTGTTTAATAAATTAGTTTGGGTTCTCTACATGCATAACAAAC 358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes an isolated polypeptide (I) comprising at least an immunogenic portion of a prostate tunnour antigen protein or its variant. (I) have cytostatic activity and can be used in vaccine production. (I), prostate tunnour antigen polynucleotides, an antigen presenting cell (Apc e.g. a dendritic cell) that expresses (I), and a pharmaceutical composition containing (I) are useful for inhibiting the specific proteins and oligonucleotides that hybridise to a
                                           TGTTGC--AAAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTT
                                                                                                                               GCTTTTTCCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAACATCTGA
                                                                                                                                                 2653 GCTTTTCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAACATCTGA
                                                                                                                                                                                                                239 AGAGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCA
                                                                                                                                                                                                                                       CCTGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCACCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            419 ACTITATITICTATGTGTTTTTTGCAACATATGAGTGTTTTGAAAATAAAGTACCCATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Prostate specific protein and its encoding polynucleotide, useful for the treatment and diagnosis of prostate cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tumour antigen; prostate tumour; therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prostate tumour antigen determined cDNA sequence for P705P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; prostate tumour antigen; prostate tumour; thera; prostate cancer; immunogenic; cytostatic; vaccine; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 50; Page 219; 276pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-OCT-2000; 2000WO-US27464.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Xu J, Skeiky YAW,
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The present invention describes an isolated polypeptide (PI) comprising at least an immunogenic portion of a prostate-specific protein, or its variant. Also described are polynucleotides (NI) encoding (PI). (PI) and The polypeptides activity and can be used in vaccine production. The polypeptides, nucleic acids and antibodies from the present Invention are useful in the diagnosis and therapy of prostate cancer. Prostate specific genes P704P, P712P, P74P, P775 and B315D are located in a genomic region on chromosome 22q11.2 known as the Cat Eye Syndrome chromosome 1. AAH84671 to AAH885143 and AAG99000 to AAG99077 represent polynucleotide and polypeptide sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9
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                                                                                                                                                                                                                                                                                                                                                                                                             Human; prostate cancer; therapy; diagnosis; cat eye syndrome; chromosome 22q11.2; prostate-specific protein; chromosome 1; prostate specific antigen; PSA; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 472; DB 22; Length 2984;
Pred. No. 3.2e-113;
0; Mismatches 0; Indels 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cham JL, Harlocker SL, Jiang Y,
Stolk JA, Day CH, Skeiky YAW,
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                                                                                                                                                                                                                                                                                                                                                                 Human prostate-specific cDNA sequence P705P/9-F3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 31; Page 233-234; 325pp; English.
                                                                                                                                                                                                                                              BP.
                                                                                                                                                                                                                                     AAH84994 standard; cDNA; 2984
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99US-0443686,
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                                                                                                                     2953 TCTTTA 2958
                                                                                        TCTTTA 484
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200134802-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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18-NOV-1999;
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                                                                                                                                                                                                                                                                                AAH84994;
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Best Local Si
Matches 484;
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                                                                                                                                                                                                                                    AGAGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCA 298
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                                                                                                           AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTG
                                                                                                                                           TGTTGC -- AAAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTT
                                                                                                                                                                           GCTTTTTCCCCCATTGGAACTAGTCATTAACCCCATCTGTAAACTTGGTAGAAAAACATCTGA
                                                                                                                                                                                                            GACAGCCTGTTTCTATCCTGTTTAATAAATTAGTTTGGGTTCTCTACATGCATAACAAAC
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polynucleotide that encodes a prostate specific protein are useful for detecting the presence or absence of a cancer or monitoring the progression the progression of a cancer, especially prostate cancer. AAH03422 to AAH3872, AAB14798 to AAB74821 and AAB74830 are sequences used in the exemplification of the present invention.
                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunotherapy; cancer; leukaemia; WT1; Wilm's tumour
v13; zinc finger transcription factor; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human P705P inventive antigen coding sequence SEQ ID NO: 349.
                                                                                   Indels
                                                                  Length
                                                  Sequence 2984 BP; 837 A; 659 C; 645 G; 842 T; 1 other;
                                                                                   ö
                                                                  Score 472; DB 22;
Pred. No. 3.2e-113;
                                                                                   Mismatches
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                                                                  97.5%;
99.6%;
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                                                                                   Conservative
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|TCTTTA 2958
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                                                                                                                                                                                                             The present invention describes compositions comprising peptides derived from the Wilm's tumour protein WT1 and methods for their use in treating malignant diseases. Peptides derived from both the murine and human WT1 proteins are provided. The human WT1 gene is found on chromosome 11p13, and the protein was shown to be a zinc finger transcription factor. The dimmunogenic peptides of the invention are particularly useful in the diagnosis and treatment of cancer and leukaemia. The present sequence is coding sequence used in the exemplification of the invention.
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AA7283 2024g07.s

AA198281 10 prosts

AA809587 nj43e11.s

AI685682 tt89h03.x

BI855676 60383205

AA229495 nc35e03.s

AA4028 nr28d08.r

AA67028 nr28d08.r

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AA533766 nj56e4.s

AA533760 nc35e04.s
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AA4041 nc35h08.s
AA550104 nc35h08.s
BF095382 1L2-UT007
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AA650230 ns93e02.s
AA572913 nm42f12.s
BI360784 388827 MA
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A1027196 ov83a09.x
AA507804 ng89g12.s
AW135064 UI-H-BI1-
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I Chases I to 538)

NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.

NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index

Onpublished

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Michael J. Brownstein, M.D., Ph.D., Michael R.
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AI829793 wc07c11.x
AA640241 nr20h08.s
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                                                                   mRNA sequence.
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Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

AI620178 tu54a11.x BQ018482 UI-H-DHI-BQ020946 UI-H-DHI-AA813266 aj44g03.s

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484 484 483 473

Description

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Query Match I

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Result . 80

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/organism="Homo sapiens"
//mol_type="mRNA" sapiens"
//mol_type="mRNA" 1960;"
//clone="InAACR: 1960;"
//dev_stage="Adult"
//dev
                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.

Email: cgapbs-rémail.nih.gov
Tissue Procurement: Dr. Jose Mercuende
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Clone distribution information can be found
through the I.M.A.G.E. Consortium/LIML at: http://mage.llnl.gov
sequence: 1-30, >AT_lich*Low_complexity (matched compliment)
POLYA-Yes.
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1 (Dases 1 to 663)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Unpublished
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                                                                                                                                                                                                                                           /dev_stage="adult"
//dev_stage="adult"
//dev_stage="adult"
//deb_host="DH10B"
//clone_lib="ACCGAP_Pr28"
//note="Organ" NCI_CGAP_Pr28"
//note="Organ" NCI_CGAP_Pr22 was prepared, and so inormalized library NCI_CGAP_Pr22 was prepared, and ss circles were made in vitro. Following HAP purification, reaction. The driver was PCR-amplified cDNAs from a pool 5,000 clones made from the same library (cloneIDs 985608-986759, 1101192-1101959, and 1217928-1220615).
Subtraction by Bento Soares and M. Fatima Bonaldo. ""
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATTG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 484; DB 9;
100.0%; Pred. No. 2.6e-90;
1ve 0; Mismatches 0;
              Std Error: 0.00
                                                                                                                                        /organism="Homo sapiens"
                     Seq primer: -40UP from Gibco
High quality sequence stop: 456
                                                                                                                                                             /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:2254844"
Insert Length: 825 Std Erro
Seg primer: -40UP from Gibco
                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                  /sex="male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      484; Conservative
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Best Local Similarity
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421 TITATITITICTATGTGTTTTTGCAACATATGAGTGTTTTGAAAATAAAGTACCCATGTC 480
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sequence tag for this library is AGATCATTGC.
TAG_LIB=UT-H-DH1
TAG_LISUE=lung
TAG_SEQ=AGATCATTCC.
1 106 c 139 g 194 t 2 others
                                                                                                                           Length 661;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/Libr at:
                                                                                                                                                       1;
                                                                                                                         99.8%; Score 483; DB 12; 99.8%; Pred. No. 4e-90;
                                                                                                                                                              0; Mismatches
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UI-H-DH1-axh-j-06-0-UI.sl NCI_CGAP_DH1 Homo sapiens cDNA clone
IMAGE:5828837 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs.remail.nib.gov
Tissue Procurement: Dr. Jose Mercuende
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Clone distribution information can be found
through the I.M.A.G.E. Consortium/LiNL at: http://image.llnl.gov
The following repetitive elements were found in this cDNA
Seq primer: M13 FORWARD
POLYA-YES.
                                                                                                                                      AGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCAGA 300
                    270 AGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTCACCAGA 211
                                                                    301 CAGCCTGTTTCTATCCTGTTTAATAATTAGTTTGGGTTCTCTACATGCATAACAAACCC 360
                                                                                      210 CAGCCTGTTTCTATCCTGTTTAATAAATTAGTTTGGGTTCTCTACATGCATACAAAAACCC 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryotani Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. I (bases 1 to 6d). NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
/db_xref="taxon:9
                                                                                                                                                                                                                                                                                                                                                                                                                                                              BQ020946
BQ020946.1 GI:19756232
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aj4403.su Soares_testis_NHT Homo sapiens cDNA clone 1393204 3', AM813266
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cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima Bonaldo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequencing Center
information can be
31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. L (bases 1 to 599)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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MCLTATY Labo
University of Iowa
University of Iowa
University of Iowa
University of Iowa Med Labs, Iowa City, IA 52242, USA
Tel: 319 356 4866
Fax: 319 356 7171
Email: paul-mccray@ulowa.edu
Tissue Procurement: Dr. M. J. Welsh, University of Iowa
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Research
Genetics (www.resgen.com) or from Open Blosystems
The following repetitive elements were found in this CDNA
Sequence: 1-21, >AT_rich#Low_complexity (matched compliment)
POLYA=VES:
                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                       1 (bases 1 to 674)

Bonaldo, M.F., Lennon, G. and Soares, M.B.

Normalization and subtraction: two approaches to facilitate gene
                                                                                                                                                                       Genome Res. 6 (9), 791-806 (1996)
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99.88;
                                                  Homo sapiens (human)
                                                                                                                                                                                                                      Contact: McCray, PB
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                                                                 sapiens
                BM983868.1
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         300 ACAGCCIGITICIAICCIGITIAAIAAAITAGITIIGGGITCICTACAIGCATAACAAACC 359
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                                                                                                                                                                                                                                                                                                                                                                                                               Length 599;
                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 982 Std Error: 0.00
Seg primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 503.
                                                                                                                                                                                                                                                                                                                                                                                                     97.7%; Score 473; DB 9;
99.8%; Pred. No. 4.9e-88;
Live 0; Mismatches 0;
                                                                              1. .599
.Corganism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="1393204"
                                                                 Location/Qualifiers
                                                                                                                                                        /sex="male"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                    484; Conservative
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                                                                                                          Score 473; DB 12;
Pred. No. 4.8e-88;
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DEFINITION

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Tissue Procurement: James Martin
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Clone distribution information can be obtained
from Dr. M. Bento Soares, bento-soares@ulowa.edu
POLYA-Yes.
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TGTTGCAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTGC 180
                                                                                                                                                                         CAGCCTGTTTCTATCCTGTTTAATAAATTAGTTTGGGTTCTCTACATGCATAACAAACCC 360
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                                                                                                                                                                                                                                                                                                                       23
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 631)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCI CGAP http://www.ncbl.nlm.nih.gov/nciogap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                031 bp mRNA linear EST 07-NOV-UI-H-FLI-bge-i-17-0-UI-BI NCI_CGAP_FLI Homo sapiens cDNA clone UI-H-FLI-bge-i-17-0-UI 3', mRNA sequence.
                                                        TTTTTCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAAAACTCTGAAG
                                                                       AGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCAGA
                                                                                                                                           262 AGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCAGA
                                                                                                                                                                                                                                361 TGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCCA-CCAAA
                                                                                                                                                                                                                                               142 TGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCACAAA
                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref-"taxon:9606"
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EST.
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                                                                                                                                                                                                                                                                                                                                                  CTTTA 484
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CA430903/c
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with an oligo-dr primer containing a Not I site. Double stranded CDNA was ligated to an ECOR I adaptor, digested with Not I, and cloned directionally into pyTy3-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dTy18 tail. The sequence tag for this library is GAGGTCGTG. The cell lines were provided by Dr. James Martin from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AMM43/224 490 bp mRNA linear EST 30-MAY-1997 2v54b10.s1 Soares_testis_NHT Homo sapiens cDNA clone IMAGE:757435 3' similar to contains Alu repetitive element;, mRNA sequence. AA437224
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Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S.,
Kucaba,T., Lecy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B.,
Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,
                                                                                                                                                                                                                                                                                                                                                                                                                                    505 TITIAAAATAAGCATITAGIGCICCAGICCCTACIGAGIACICITICICICCCCCCCCTCTG 446
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  University of Iowa.
TAG_LIB=UI-H-FL1
TAG_TISSUE=Human Chondrosarcoma Grade 3 cell line mix
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTG
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                                                                                                                                                                                                                                                                                                               Score 472; DB 14;
Pred. No. 7.8e-88;
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102 c 136 q
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Similarity
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                                            RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTGC---AAAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTG 179
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                                                 Tel: 34 286 1800
Fax: 314 286 1800
Fax: 314 286 1810
Fax: 314 286 1810
Final clore is available royalty-free through LLNL; contact the IMAGE Consortium (infe@image.llnl.gov) for further information. Seq primer: -41m13 fwd. ET from Amersham
High quality sequence stop: 363.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTTTATTTTTCTATGTGTTTTTTGCAACATATGAGTGTTTTGAAAATAAAGTACCCATGT 479
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                               Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                            Score 469; DB 9; Length 490;
Pred. No. 3.4e-87;
0; Mismatches 0; Indels
                                                                                                                                       1. 490
/organism="Homo sapiens"
/mol_type="mrnn"
/db_xref="GDB:5978049"
/db_xref="taxon:9606"
                                                                                                                                                                                             /clone="IMAGE:757435"
  WashU-Merck EST Project 1997
                                                                                                                                                                                                                 /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                           tch 96.9%;
al Similarity 99.4%;
482; Conservative
                                                                                                                                                                                                        /sex="male"
                      Contact: Wilson RK
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                                                                                                                                                                                                                                                                                                                                            173
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//db_xref="taxon:9166"
/db_xref="taxon:97820"
/db_host="NAGE:1577820"
/lab_host="NH10B"
/clone_lib="soares_NFL_T_GBC_SI"
/note="Organ: pooled; Vector: p7773D-Pac (Pharmacia) with a modified polylinker; Site_l: Not I; Site_2: Ecc RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCBI) were mixed, and as circles were made in vitro. Pollowing HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver from the same 3 libraries. The pools consisted of I.M. A.G.E. clones 297480-302897, 682632-687239, Soares and M. Fatima Bonaldo."
                  EST 07-JUL-1998
                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 455)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (inf@image.llnl.gov) for further information.
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 443.
Location/Qualifiers

1. 455
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AA9/2883 AOD AA9/2883 AOD AA9/2883 AOD AA9/2883 AOD AA9/2889 AOD AA9/2
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100.0%; Pred. No. 1.3e-82;
ive 0; Mismatches 0;
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/mol_type="mRNA"
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Homo sapiens
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Contact: Robert Strausberg, Ph.D.

Email: cgapbs-rémail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: David B. Krizman, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center-
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.linl.gov/bbrp/image/thage.html
Insert Length: 605 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham.

Location/Qualifiers

rce
                                                                                                                                                                                                                                                                                                     AA809587 435 bp mRNA linear EST 18-FEB-1998 nj43ell.sl NCI_CGAP_Pr9 Homo saplens cDNA clone IMAGE:995276, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="normal prostatic epithelial cells"
/lab_host="DH10B"
/lab_host="DH10B"
/clone_lib="NCI_CGAP_FP9"
/note="Organ: prostate; Vector: pAMP10; mRNA made from normal prostatic epithelial cells, cDNA made by oligo-dT priming. Non-directionally cloned. Slze-selected on agarose gel, average insert size 600 bp. Library made by D. Krizman, NIH.

a 96 c 68 g 151 t
                                                                  Craniata; Vertebrata; Euteleostomi;
Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria: Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 435)
NCI-CGAP http://www.ncbl.nlm.nlh.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index
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                                                                                                                 400 TAGICAGCACCCCACCAAACTITATITITCTATGTGTTTTTTGCAACATATGAGTGTTT
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Pred. No. 2.2e-79;
); Mismatches 1; Indels 0;
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/mol_type="mRNA"
/mb_xref="taxon:9606"
/clone="TMAGE:995276"
/sex="male"
                                                                                                                                                                                                                                                                                                                                      Sequence.
AA809587
AA809587.1 GI:2878993
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99.8%;
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LNCaP Homo sapiens CDNA, mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="mkNa"
/db_xref="taxon:9606"
/db_xref="taxon:9606"
/cell_line="LNCap"
/clole_lib="prostate cancer cell line LNCap"
/note="prostate cancer metastasized to lymph node;
subtracted cDNA libraries from prostate cancer cell line
LNCap treated with androgen."
                                                96
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                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 434)
Zhang,J.S. and Smith,D.I.
High throughput screening for androgen regulated genes in LNCaP
cells: Identification of hAG-2 as an androgen regulated gene over
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               TTCTCTACATGCATAACAAACCCTGCTCCAATCTGTCACATAAAAGTCTGTGAAG
                                 398 TTTAGTCAGCACCCCCACCAAACTTTATTTTTCTATGTGTTTTTTGCAACATATGAGTGT
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Catarrhini; Hominidae;
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100.0%; Pred. No. 6.3e-80;
.1ve 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expressed in prostate adenocarcinoma Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Zhang, Jin-San
Dept. Pathology and Lab Medicine
Mayo Clinic Cancer Center
200 lst St. SW, Rochester MN 55905, 1
Tel: 507-2660311
Fax: 507-2665193
                                                                                                                                               /organism="Homo sapiens"
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Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                     989281 434 prostate cancer cell line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: zhang,jinsan@mayo.edu.
Location/Qualifiers
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EST.
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Subtraction by Bento Soares and M. Fatima Bonaldo. 66 c \, 83 g \, 124 t
                                                                                  87.2%; Score 422; DB 9; 1
100.0%; Pred. No. 1.9e-77;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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/mol_type="mRNA"
/db_xref="taxon:9606"
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BI855676.1 GI:15996423
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Email: cgapbs-rémail.nih.gov
Tissue Procurement: Michael J. Brownstein, M.D., Ph.D., Michael R.

Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-GGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
Seq primer: -400P from Gibco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI685682 423 bp mRNA linear EST 27-MAY-1999 ML4B9h03.x1 NCI_CGAP_Pr28 Homo sapiens cDNA clone IMAGE:2248757 3', AI685682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="Organ: Prostate; Vector: pT773D-pac (Pharmacia)
with a modified polylinker; plasmid DNA from the
normalized library NCL_CGAP_pr22 was prepared, and ss
circles were made in vitro. Following HAP purification,
this DNA was used as tracer in a subtractive hybridization
reaction. The driver was PCR-amplified DNAs from a pool
of 5,000 clones made from the same library (cloners)
985608-986759, 1101192-1101959, and 1217928-1220615).
201
                                                  123 TCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTGCTTTTTCCCCATTGGAACTAGT 182
                                                                                                                            322 AATAAATTAGTTTGGGTTCTCTACATGCATAACAAACCCTGCTCCAATCTGTCACATAAA 381
                                                                                                                                                                                                                                                                                                362
                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammaliai; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 423)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NAtional Cancer Institute, Cancer Genome Anatomy Project (CGAP), Unpublished
                                   TCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTGCTTTTTCCCCATTGGAACTAGT
                                                                                                                                                                                                                                                                       202 CATTAACCCATCTCTGAACTGGTAGAAAACATCTGAAGAGCTAGTCTATCAGCATCTGA
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone='IMAGE:2248757"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /dev_stage="adult"
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                    /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                              442 TGCAACATATGAG 454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .423
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DEFINITION
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AI685682/c
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63 TITAATICITICAACITGCAATTIGCAAGGATTACACATTICACIGIGAIGTATATIGIG 122
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                                                                                               123 TTGCAAAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTGCTT 182
                                                                                                                                                                                                                                                                                                                                 302
                                                                                                                                                                             362
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1 (bases 1 to 607)

NIH-MGC http://mgc.nci.nih.gov/.

National institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                        243 CTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTCACCCAGACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            123 CTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCACCAACTT
                                                                                                                                                                                                                                        183 TITCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAACATCTGAAGAG
                                         Gaps
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Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM1999 row: i column: 02
High quality sequence stop: 586.
Location/Qualifiers
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                                           ö
Length 423;
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/olone_lib="NCI_CGAP_Pr2"
/olone_lib="NCI_CGAP_Pr2"
/note="Vector: pAMP10; Site_1: Not1; Site_2: EccRI; 1st
strand cDNA was primed with oligo(dT)17 on 50 ng of
DNAse-treated, total cellular RNA obtained from 5,000-10
,000 microdissected preneoplastic cells
histologically-determined to be prostatic intraepithelial
neoplasia 2 (PICJ) cells. Double-stranded cDNA was
ligated to EccRI adaptors, 5 cycles of PCR applied to the
cDNA with an adaptor-specific primer, and the resulting
PCR product subcloned into pAMP10 by the UDG-cloning
method (Life Technologies). Average insert size is 600
bp. NOTE: Not directionally cloned. This library was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 411 bp mRNA linear EST 27-OCT-1997
NCI_CGAP_Pr3 Homo sapiens cDNA clone IMAGE:1169295, mRNA
Clone distribution: NCI-CGAP clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
www.bio.llnl.gov/bbrp/image/image.html
Insert Length: 856 Std Error: 0.00
Seg primer: -4,11n3 fwd. ET from Amersham
High quality sequence stop: 368.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 386.6; DB.9;
Pred. No. 4.3e-70;
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9 q 131 t
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                                                                                                                                                                                                                           /sex="Male"
/dev_stage="45 years old"
/lab_host="DH10B"
                                                                                                                                                 /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                            /clone="IMAGE:1010092"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         by 79 g
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Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.
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                   /tissue_type="mammary adenocarcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_BR="
/clone_lib="NIH_MGC_BR="
/clone_lib="NIH_MGC_BR="
/clone_lib="NIH_MGC_BR="
/clone_lib="NIH_MGC_BR="
/clone_lib="Nite_lib"
/clone_lib="Noti;
/cloned unidirectionally; oligo-dr primed.
/clone_lib lib lib library enriched for average insert size 1.383 kb. Library enriched for full-length clones and constructed by Life Technologies.
/clone lib lib library."
/clone_lib library."
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NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Center
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COMA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon,
DNA Sequencing by: Washington University Genome Sequencing
                                                                                                                                                                                                                                                                  9;
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                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                10;
                                                                                                                                                                                                                         Score 389; DB 12;
Pred. No. 1.3e-70;
0; Mismatches 10;
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Contact: Robert Strausberg, Ph.D.
     /clone="IMAGE:5392201"
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                                                                                                                                                                                                                           Query Match 80.4%;
Best Local Similarity 96.6%;
Matches 451; Conservative
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197 300

Gaps

240

257

180

377

437 9

EST 18-DEC-1999

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AI672753

383 bp mRNA linear EST 18-DEC-1999; we58010.x1 Soares_thymus_NHFTh Homo sapiens cDNA clone IMAGE:2345299 3' similar to contains Alu repetitive element;, mRNA
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I (bases 1 to 383)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library constructed by Bento Soares and M. Fatima Bonaldo. " 63 c 75 g 111 t
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This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 1465 Std Error: 0.00 Seq primer: -400P from gibco High quality sequence stop: 362.
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Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                    Contect: Robert Strausberg, Ph.D.
Email: gapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.
ONA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Parayed by: Genome Systems Inc., Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
found through the I.M.A.G.E. Consortium/LINL at:
Seq primer: -28m13 revl ET from Amersham
High quality sequence stop: 387.

Location/Qualifiers

Location/Qualifiers
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                                                                     1 (bases 1 to 411)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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APPLICANT: Xi, Jiangchun C. APPLICANT: Dillou, Davin C. APPLICANT: Dillou, Davin C. APPLICANT: Dillou, Davin C. APPLICANT: Michael Cart. Sean Louise APPLICANT: Harloam, Janay Yuqui APPLICANT: Harloam, Salvan G. APPLICANT: Harloam, Salvan G. APPLICANT: Raios, Michael APPLICANT: Raios, Michael APPLICANT: Bay Carig APPLICANT: Bay Carig APPLICANT: Bay Carig APPLICANT: Bay Carig APPLICANT: DAGNOSIS OF PROSTATE CANCER TITLE OF INVENTIONS: DAGNOSIS OF PROSTATE CANCER TITLE OF INVENTIONS: DAGNOSIS OF PROSTATE CANCER TITLE OF INVENTIONS: DAGNOSIS OF PROSTATE CURRENT APPLICATION NUMBER: Usy09/439,313 CURRENT APPLICATION NUMBER: Usy09/439,313 CURRENT APPLICANTON NUMBER: Usy09/439,313 CURRENT APPLICANTON NUMBER: Usy09/439,313 OURING OF SEQ ID NOS: 575 SEQ ID NO 434 CREAKISM: Homo sapiens 100.0%; Fored. No. 6.5e-121; Matches 484; Conservative 0; Mismatches 0; Indels 0; Gaps 1 TITTAAAAATAAATAAATAACCATTAGCACTCACACACACAC		atent No. 63295Us Eneral, information:
APPLICANT: Dillon_Davin C. APPLICANT: Harlocker, Jennifer L. APPLICANT: Harlocker, Jennifer L. APPLICANT: Jiang Yugui APPLICANT: Jiang Yugui APPLICANT: Rados, Michael APPLICANT: Rater, Mark APPLICANT: Rater, Mark APPLICANT: Solk, John APPLICANT: Solk, John APPLICANT: Day CURRENT FILING DATE: 1999-11-12 CURRENT FILING DATE: 1999-11-12 CURRENT FILING DATE: 1999-11-12 SOFTWARE: PRASED for Windows Version 3.0 CURRENT FILING DATE: 1999-11-12 SOFTWARE: PRASED for Windows Version 3.0 CURRENT FILING DATE: 1999-11-12 SOFTWARE: PRASED for Windows Version 3.0 CURRENT FILING DATE: 1999-11-12 SOFTWARE: PRASED for Windows Version 3.0 CURRENT FILING DATE: 1999-11-12 SOFTWARE: PRASED for Windows Version 3.0 CURRENT H. 484 TYPE: DNA TYPE DAY	·	APPLICANT: Xu, Jiangchun
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APPLICANY: Jiang Yugui APPLICANY: Read, Steven G. APPLICANY: Panger, Gary APPLICANY: Panger, Gary APPLICANY: Panger, Gary APPLICANY: Day, Craig TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REPERREC: 120121, 47709 CURRENY FILLING DAFE: 1999-11-12 NUMBER OF EQ. DAYOS: 575 CURRENY FILLING DAFE: 1999-11-12 NUMBER OF EQ. DAYOS: 575 CURRENY FAST AND ALCANAMISH: Homo sapiens OBST LOCAL AND SAME AND ARACHISM: Homo sapiens OBST LOCAL AND SAME AND ORANISM: Homo sapiens OBST LOCAL AND SAME AND ORANISM: HOMO SAPIENS OBST LOCAL AND SAME AND ARACHISM: HOMO SAPIENS OBST LOCAL AND SAME AND ARACHISM: HOMO SAPIENS OBST LOCAL AND SAME AND ARACHISM: HOMO SAPIENS OBST LOCAL AND SAME		
APPLICANT: Reed, Steven G. APPLICANT: Read, Steven G. APPLICANT: Ralos, Michael APPLICANT: Fanger, Gary APPLICANT: Fanger, Gary APPLICANT: Fanger, Gary APPLICANT: Fanger, Gary APPLICANT: Solk, John APPLICANT: Solk, John APPLICANT: Day, Craig TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.427C9 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FASTSOR FOR VERSION 3.0 SEQ ID NO 434 LENGTH: 484 LENGTH: 484 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 CURRENT FASTSOR FOR THE CANADA CONTENT CO		
APPLICANT: Falos, Michael APPLICANT: Falos, Gary APPLICANT: Retter, Mark APPLICANT: Retter, Mark APPLICANT: Retter, Mark APPLICANT: Retter, Mark APPLICANT: Solk, John APPLICANT: Bolk, John APPLICANT: Day, Craig TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REPRENCE: 1999-11-12 CURRENT APPLICATION NUMBER: US/09/439,313 CURRENT ELLING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 SOFTWARE: FEASTSQ for Windows Version 3.0 SEQ ID NO 434 TYPE: DAA TYPE: DAA ONGANISH: Homo sapiens 100.0%; Pred. No. 6.5e-121; Best Local Similarity 100.0%; Pred. No. 6.5e-121; Best Local Similarity 00.0%; Pred. No. 6.5e-121; Best Local Similarity 1101.0%; Pred. No. 6.5e-121; Batches 484; Conservative 0; Mismatches 0; Indels 0; Gaps 1 TTTAAAATAATAAGCATTAGGGCTCAGTCCCTACGGTACTTTCACTGGTAGAATGTATATTG 12 ITTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	•••	
APPLICANT: Retter, Gary APPLICANT: Retter, Gary APPLICANT: Banger, Gary APPLICANT: Bolk, John APPLICANT: Solk, John APPLICANT: Solk, John APPLICANT: Solk, John APPLICANT: Solk, John APPLICANT: Day, Craig TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: COMPOSITIONS OF PROSTATE CANCER FILE REFERENCE: 210.121.427C9 CURRENT APPLICATION NUMBER: US/09/439,313 CURRENT APPLICATION NUMBER: US/09/439,313 NUMBER OF SEQ ID NOS: 575 SOFTWARE: FastSEQ for Windows Version 3.0 SOFTWARE: FastSEQ for Windows Version 3.0 SOFTWARE: PASTSEQ for Mindows Version 3.0 ORGANISM: Homo sapiens ONGANISM: Homo sapiens 100.0%; Score 484; DB 4; Length 484; Duery Match Best Local Similarity 100.0%; Pred. No. 6.5e-121; CONSETVATIVE OF MISSMACCATTACTCCCCCTCCTCTCG 60 ITTTAAAATAAATAACCATTACGCACTCCCACTCCACTCACT		
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APPLICANT: Day, Crait APPLICANT: Day, Crait APPLICANT: Day, Crait TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: COMPOSITION: COMPO		
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.27C9 CURRENT FILING DATE: 1999-11-12 NUMBER OF SEQ ID NOS: 575 SOFTWARE: FastSEQ for Windows Version 3.0 SO	٠.	
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER CURRENT AND STATE STATE CANCER CURRENT APPLICATION NUMBER: 1999-11-12 CURRENT FILING DATE: 1999-11-12 CURRENT FILING DATE: 1999-11-12 SEQ ID NO 434 LENGTH: 484 LENGTH: 484 TYPE: DNA ORGANISM: Homo sapiens 09-439-313-434 ORGANISM: Homo sapiens 100.0%; Score 484; DB 4; Length 484; CORSERVATIVE DNA ORGANISM: Homo sapiens 1 TTTAAAATAAGCATTTAGTGCTCAGTCCTACTGGAGTACTTCTCTCCCCTCTG 60 1 TTTAAAATAAGCATTTAGTGCTCAGTCCTACTGGAGTACTTTCTCTCCCCTCTTG 60 1 TTTAAAATAAGCATTTAGTGCTCAGTCCTACTGGAGTACTTTCTCTCTC		
FILE REFERENCE: 210121.427C9  CURRENT APPLICATION NUMBER: US/09/439,313  CURRENT FILING DATE: 1999-11-12  UNUMBER OF SEQ ID NOS: 575  SOFTWARE: FastSEQ for Windows Version 3.0  SEQ ID NO 434  LENGTH: 484  TYPE: DN 434  LENGTH: 484  TYPE: DN 63; SCORE 484; DB 4; Length 484;  COBAISM: Homo sapiens  100.0%; SCORE 484; DB 4; Length 484;  ONGANISM: Homo sapiens  101-17-17-17-17-17-17-17-17-17-17-17-17-17		
CURRENT FILING DATE: 1999-11-12  NUMBER OF SEQ ID NOS: 575  LENGTH: 484  LINDO 084; SCORE 484; DB 4; Length 484;  ORGANISM: Homo sapiens  ORGANIS		FILE REFERENCE: 210121,427C9
NUMBER OF SEQ ID NOS: 575  SECTION NO. 1434  LENGTH: 484  LENGTH: 484  LENGTH: 484  LENGTH: 484  LENGTH: 484  LENGTH: 484  ORGANISM: Homo sapiens  100.0%; Score 484; DB 4; Length 484;  ORGANISM: Homo sapiens  100.0%; Pred. No. 6.5e-121;  Matches 484; Conservative 0; Mismatches 0; Indels 0; Gaps  1 TTTAAAATAAGCATTTAGTGCTCAGTCCTAGTAGTACTTTCTCCCCTCTG 60		CURKENT APPLICATION NUMBER: 05/09/439,313
SOFTWARE: FastSEQ for Windows Version 3.0  SOFTWARE: FastSEQ for Windows Version 3.0  LENGTH: 484  TYPE: DNA  ORGANISM: Homo sapiens  ORGANISM: Homo sapiens  100.0%; Score 484; DB 4; Length 484;  Duery Match  100.0%; Pred. No. 6.5e-121;  ATTTAAAATAAGCATTTAGTGCTCAGTCCTAGTAGTTCTTCTCCCCTCTG 60		NIMBER OF SET IN NOS: 575
SEQ ID NO 434		stSEQ for Windows Version 3
LENGTH: 484  TYPE: DNA ORGANISM: Homo sapiens -09-439-313-434  Duery Match Best Local Similarity 100.0%; Score 484; DB 4; Length 484;  Matches 484; Conservative 0; Mismatches 0; Indels 0; Gaps  1 TITTAAAATAAGCATTAGTGCTCACTCCTACTGAGTACTCTCTCCCCTCTG 60   IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	••	•
TYPE: DB 4; Length 484;  OKGANISM: Homo sapiens  100.0%; Score 484; DB 4; Length 484;  Duery Match  Best Local Similarity 100.0%; Pred. No. 6.5e-121;  Matches 484; Conservative 0; Mismatches 0; Indels 0; Gaps  1 TTTAAAATAAGCATTTAGTGCTCACTCGTAGTACTCTTCTCCCCTCTG 60  1 ITTTAAAATAAGCATTTAGTGCTCACTCGTAGTACTCTTTCTCCCCTCTTG 1  1 TTTTAAAATAAGCATTTAGTGCTCAAGGATTACACATTCACTGTGATGTATTG 12  1 TTTTAAAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTCACTGTGATGTATTG 12  1 GATTGCAAAAAAAAAAAAATTGCAAGTTTAAAATTACTTGGTTGTGAATTTG 12  1 TGTTGCAAAAAAAAAAAAAAATTGCAAGTTTAAAATTACTTGGTTGTGAATCTTGC 18  1 TTTTTCCCCATTGGAACTTGTTTAAATTACTTGGTTGTGAATCCATCTTGC 18  1 TTTTTCCCCATTGGAACTAGTCTTTAAAATTACTTGGTTGTGAAAACATCTGC 18  1 TTTTTCCCCATTGGAACTAGTCATTAAACTTGCTTGGTAGAAAAAAAA	••	LENGTH: 484
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Best Local Similarity 100.0%; Pred. No. 6.5e-121; Matches 484; Conservative 0; Mismatches 0; Indels 0; Gaps  1 TITAAAATAAGCATITAGTGCTCAGTCCCTAGTAGTACTTTCTCTCCCCTCTG 60  1 ITITAAAATAAGCATITAGTGCTCAGTCCTAGTAGTACTTTCTCTCCCCTCTG 60  61 AATTTAAATCTTTCAACTTGCAAATTTGCAAGGATACAATTCACTGTAATTG 12	ŭ,	100.0%; Score 484; DB 4;
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	· vo	AATTTAATTCTTTCAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTG
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181 TTTTCCCCAFTGGAACTAGTCAFTAACCCATCTCTGAACTAGAAAAACATCTGAAG 241 AGCTAGTCATCAGCAGTGACAGTGGAATTGGATGGTTCTCAGAACATTTCACCCAA	QY	TTTTCCCCCATTGGAACTAGTCATTAACCCCATCTCTGAACTGGTAGAAAAAAATTCTGAAG
241 AGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCAGA	qq	TITITCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAAAACATCTGAAG
	Qy	AGCTAGTCTATCAGCATCTGACAGGTGAATTGGATGGTTCTCAGAACCATTTCACCCAGA

Db 361 TGCTCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTTTAGTCAGCACCCCCACCAAC 420  Qy 421 TTTATTTTCTATGTGTTTTTTGCAACATATGAGTGTTTTGAAAATAAAGTACCATGTC 480	CURRENT APPLICATION NUMBER: US/09/328,475C CURRENT FILING DATE: 1999-06-09 NUMBER OF SEQ ID NOS: 341 SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 334 LENGTH: 2051 TYPE: DNA ORGANISM: Homo sapien FEATURE: LOCATION: (1)(2051) OTHER INFORMATION: (1)(2051)	Dest Local Similarity   97.3%;   Score 4/3;   Dest Length 2051;     Matches
Db 241 AGCTAGTTATATATATATATATAGTTTGATCACCAGA 300  Qy 301 CAGCCTGTTTCTATCCTGTTTAATAATTTGGTTCTTCTACACCAACCA	; APPLICANT: Xu, Jiangchun; ALPLICANT: Xu, Jiangchun; APPLICANT: Mitchem, Jennifer Lynn TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE CURRENT APPLICATION NUMBER: US/09/352,616A CURRENT APPLICATION NUMBER: US/09/352,616A NUMBER OF SEQ ID NOS: 472 SOFTWARE: FastSEQ for Windows Version 3.0 SOFTWARE: FastSEQ for Windows Version 3.0 TYPE: DNA TYPE: DNA  ORCANISM: Homo sapiens US-09-352-616A-434	Query Match         100.0%;         Score 484;         DB 4;         Length 484;           Beet Local Similarity         100.0%;         Pred. No. 6.5e-121;         0;         daps         0;           Matches         484;         Conservative         0;         Mismatches         0;         Indels         0;           QY         1 TTTTAAAATAACTAGCATTAGTGCTCAGTCCTACTAGTACTCTTCTCTCCCTCC

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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jannifer L.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Harlocker, Susan Louise
APPLICANT: Rado, Michael
APPLICANT: Rados, Michael
APPLICANT: Fanger, Gary
APPLICANT: Reter, Mark
APPLICANT: CANGOS, Michael
APPLICANT: CANGOS, Wichael
APPLICANT: CANGOS, Wichael
APPLICANT: Day, Craig
TILLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TILLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.42709
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE FEASTER FOR Mindows Version 3.0
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Pred. No. 1.9e-117;
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                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                      Sequence 335, Application US/09352616A

Patent No. 6395278

GENERAL INFORMATION:
APPLICANT: Dillon, Davin C.
APPLICANT: Jiang, Yuqui
APPLICANT: Jiang, Yuqui
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPUNDS FOR IMMUNOTHERAPY AND CURRENT APPLICATION NUMBER: US/09/352,616A

CURRENT FILING DATE: 1999-07-13

NUMBER OF SEQ ID NOS: 472
                                                                                                                                                                                                                                                                                                                                                                                                                     5;
                                                                                                                                                                                                                                                                                                                                                                                      Length 2984;
                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                  Score 472; DB 4;
Pred. No. 1.9e-117;
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                                                                                                                                                                                                                                                                   FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                      97.58;
99.68;
                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 99.6
Matches 484; Conservative
                                                                                                                                                                                                                                                                                                                                   ; ORGANISM: Homo sapien
US-09-352-616A-335
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RESULT 5
US-09-352-616A-335
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                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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NAME/KEY: misc_feature
                                                                                                        TYPE: DNA
ORGANISM: Homo Sapien
                                                                                SEQ ID NO 102
LENGTH: 1020
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APPLICANT:
APPLICANT:
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                APPLICANT: XV. Jiangchun
APPLICANT: W. Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE
FILE REFERENCE: 210121.4276
CURRENT APPLICATION NUMBER: 1999-01-15
NUMBER OF SEQ ID NOS: 338
SOFTWARE: FastSEQ for Windows Version 3.0
LENGH: 2984
                                                                                                                                                                                                                         Length 2984;
                                                                                                                                                                                                                                               0; Indels
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APPLICANT: Zhang, Jiamy
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll III, Eddie
APPLICANT: Endege, Wilson O.
APPLICANT: Ford, Donna M.
APPLICANT: Schlegel, Robert
APPLICANT: Schlegel, Robert
APPLICANT: Schlegel, Robert
APPLICANT: Steinmann, Kathleen E.
TITLE OF INVENTION: GENES AND GENE EXPRESSION PRODUCTS THAT
                                                                                                                                                                                                                  97.5%; Score 472; DB 4; 1
99.6%; Pred. No. 1.9e-117;
live 0; Mismatches 0;
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                                                                                                                                                                                                               Query Match
Best Local Similarity 99.6
Matches 484; Conservative
                                                                                                                                                                              ORGANISM: Homo sapien
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| TCTTTA 2958
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           GENERAL INFORMATION:
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Patent No.
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TITLE OF INVENTION: ARE DIFFERENTIALLY REGULATED IN PROSTATE CANCER FILE REFERENCE: 1532.002/200130_463
CURRENT APPLICATION NUMBER: US/09/328,475C
CURRENT FILING DATE: 1999-06-09
NUMBER OF SEQ ID NOS: 341
SOFTHARE: FastSEQ for Windows Version 3.0
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APPLICANT: Ford, Donna M.
APPLICANT: Ford, Donna M.
APPLICANT: Schlagel, Robert
APPLICANT: Schlagel, Robert
APPLICANT: Steinmann, Rathleen E.
TITLE OF INVENTION: GRESS AND GENE EXPRESSION PRODUCTS THAT
TITLE OF INVENTION: ARE DIFFERENTIALLY REGULATED IN PROSTATE CANCER
FILLE REFERENTE : 1532.002/200130.463
CURRENT APPLICATION NUMBER: US/09/328,475C
CURRENT APPLICATION NUMBER: US/09/328,475C
CURRENT FILING DATE: 1999-06-09
NUMBER OF SEQ ID NOS: 341
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                           2;
                                                                                                                                                                                                                                                                                                                                   Length 1020;
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                                                                                                                                                                                                                                                                                                                          Score 426.2; DB 4
Pred. No. 2.6e-105
0; Mismatches 18
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8-09-38-475C-103/c
5-09-38-475C-103/c
Sequence 103, Application US/09328475C
Patent No. 6476207
GENERAL INFORMATION:
                                                                                                                                                                                                                                          ; LOCATION: (1)...(1020)
; OTHER INFORMATION: n = A,T,C or G
US-09-328-475C-102
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95.78;
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APPLICANT: Astel, Jon H.
APPLICANT: Carroll III, Eddie
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Matches 449; Conservative
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; LOCATION: (1)...(822)
; OTHER INFORMATION: n = A,T,C or
US-09-328-475C-223
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                                                                                                                                                                             65
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APPLICANT: Astel, Jon H.
APPLICANT: Astel, Jon H.
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll III, Eddie
APPLICANT: Ford, Donna M.
APPLICANT: Ford, Donna M.
APPLICANT: Schlegel, Robert
APPLICANT: Schlegel, Robert
TITLE OF INVENTION: GREE AND GENE EXPRESSION PRODUCTS THAT
TITLE OF INVENTION: ARE DIFFERENTALLY REGULATED IN PROSTATE CANCER
FILE REFERENCE: 1532.002/200130.463
CURRENT FILING DATE: 1999-06-09
NUMBER OF SEQ ID NOS: 341
SSQ ID NO 223
LENGTH: 822
                                                                                                                                                                           6 AAATAAGCATTTAGTGCTCAGTCCCTACTGAGTACTCTTTCTCTCCCCTCCTCTGAATTT
                                                                                                                                                                                           CAAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTGCTTTTT
                                                                                                                                                                                                                                                                                                      C--AAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTGCTTTTT
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                                                                                                                           Score 426.2; DB 4;
Pred. No. 2.6e-105;
                                                                                                                                                   0; Mismatches
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Patent No. 6476207
GENERAL INFORMATION:
                       TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
NAME/KEY: misc_feature
LCCATION: (1)...(1021)
CTHER INFORMATION: n = A,T,C or G
                                                                                                                           88.18;
95.78;
                                                                                                                                       Best Local Similarity 95.7
Matches 449; Conservative
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ORGANISM: Homo Sapien
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SEQ ID NO 103
LENGTH: 1021
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                                                                                                                                                                                                                                                                                                                         GTTCTCAGAACCATTTCACCCAGACAGCCTGTTTCTATCCTGTTTAATAATTAGTTTGG
                                                                                                                                                                                                                                                                                                                                                                                                  510 GTACTCTTTCTCCCCCCCCCTCTGAATTTAATTCTTTCAACTTGCAATTTGCAAGGATTA
                                                                                                                                                     ACTTGGTTTGTGAATCCATCTTGCTTTTTCCCCATTGGAACTAGTCATTAACCCCATCTCT
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                                                                            GTACTCTTTCTCCCCCCCCCCCCCTCGAATTTAATTCTTTCAACTTGCAATTTGCAAGGATTA
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TITLE OF INVENTION: GENES AND GENE EXPRESSION PRODUCTS THAT
TITLE OF INVENTION: ARE DIFFERENTALLY REGULATED IN PROSTATE CANCER
FILE REFERENCE: 1532.002/200130.463
CURRENT FILING DATE: 1999-06-09
NUMBER OF SEQ ID NOS: 341
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 222
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   Length 822;
                                       Indels
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Score 415.6; DB 4;
Pred. No. 1.7e-102;
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Pred. No. 9e-94;
0; Mismatches
                 Pred. No. 1.76
0; Mismatches
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APPLICANT: Zhang, Jimmy
APPLICANT: Carrol, Jon H.
APPLICANT: Carroll III, Eddie
APPLICANT: Endege, Wilson O.
 85.9%;
98.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 79.1%;
Best Local Similarity 98.8%;
Matches 397; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Carroll III, Eddie
APPLICANT: Endege, Wilson O.
APPLICANT: Ford, Donna M.
APPLICANT: Monahan, John E.
APPLICANT: Schlegel, Robert
APPLICANT: Schlegel, Robert
Query Match 85.9 Best Local Similarity 98.4 Matches 431; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Homo Sapien
US-09-328-475C-222
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	Qy 40 CTCTTTCTCTCCCCCTCTGAATTTAATTCTTTCAACTTGCAAGGATTACAC 99			TC 404      TC 366	RESULT 12 US-09-030-607-115 Sequence 115, Application US/09030607 Patent No. 6262245 CENERAL INFORMATION	APPLICANT: Xu, Jiangchun APPLICANT: Xu, Jiangchun TITLE OF INVENTION: COMPGUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS NUMBER OF SEQUENCES: 224	CURKESPONDENCE ADDRESS:  ADDRESSE: SEED and BERRY LLP  STREET: 6300 Columbia Center, 701 Fifth Avenue  CITY: Seattle	; STATE: WA ; COUNTRY: USA ; ZIP: 98104 ; COMPUTER READABLE FORM:		CURRENT APPLICATION DATA:  APPLICATION NUMBER: US/09/030,607  FILING DATE: 25-FFR-1948	CLASSIFICATION:  ATTORNEY/AGENT INFORMATION:  NAME: MAKI, David J.	TELEPHONE: (206) 622-4900	; TELEFAX: (206) 682-6031 ; INFORMATION FOR SEQ ID NO: 115: ; SEQUENCE CHARACLERISTICS: ; TARNITH: 366 base continued.	TYPE: nucleic acid STRANDEDNESS: single	# MOLECULE TYPE: CDNA # ORGANISM: Homo Sapiens US-09-030-607-115
66	OY 97 CACATTICACTGTGTATATTGTGTTGCAAAAAAAAAGGTGTCTTTGTTAAAATT 156 			RESULT 11 US-09-020-956-115 : Sequence 115, Application US/09020956 ; Patent No. 6261562	GENERAL INFORMATION:  APPLICANT: Xu, Jiangchun  TTYLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS FO  NUMBER OF SEQUENCES: 178	CORRESPONDENCE ADDRESS:  STREET: 6300 Columbia Center, 701 Fifth Avenue  STATE: Asattle	COUNTRY: USA ZIP: 98104 ; COMPUTER READABLE FORM:	HELLOW TYPE: Floppy disk COMPUTER: IBW PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOSTWARE: Patentin Release #1.0, Version #1.30	9/020,956	; ATTORNEY/AGENT INFORMATION: ; NAME: Maki, David J. ; REGISTRATION NUMBER: 31,392	KEFERENCE/DOCKET NUMBER: 210121.427C2- FELECOMMUNICATION INFORMATION: FELEPHONE: (206) 522-4900	; INFORMATION FOR SEQUENCE CHARACTERISTICS: ; LENGTH: 366 base pairs	/ TYPE: nucleic acid / STRANDEDNESS: single // TOPOLOGY: linear // MOLECULE TYPE: CDNA	; ORIGINAL SOURCE; ; ORGANISM: Homo Sapiens US-09-020-956-115	Query Match 75.4%; Score 365; DB 3; Length 366; Best Local Similarity 100.0%; Pred. No. 4.9e-89; Matches 365; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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100 CICTACATGCATAACAAACCCTGCTCCCAATCTGTCACATAAAAGTCTGTGACTTGAAGTT
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APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Harlocker, Susan Louise
APPLICANT: Harlocker, Susan Louise
APPLICANT: Attochem, Jennifer Lynn
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE REFERENCE: 210121.427C8
CURRENT APPLICATION NUMBER: US/09/352,616A
CURRENT FILING DATE: 1999-07-13
NUMBER OF SEQ ID NOS: 472
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US-09-352-616A-115
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Best Local Similarity
Matches 365; Conserv
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US-09-352-616A-115
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APPLICANT: Xu, Jiangchun

APPLICANT: Dillon, Davin C.

APPLICANT: Mitcham, Jennifer L.

APPLICANT: Mitcham, Jennifer L.

APPLICANT: Harlocker, Susan Louise

APPLICANT: Reed, Steven G.

APPLICANT: Reados, Michael

APPLICANT: Reter, Mark

APPLICANT: Retter, Mark

APPLICANT: Retter, Mark

APPLICANT: Solk, John

APPLICANT: OINVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

FILE REFERENCE: 210121.427C9

CURRENT FILING DATE: 1999-11-12

CURRENT FILING DATE: 1999-11-12

SOFTWARE: FastSEQ for Windows Version 3.0

SOFTWARE: FastSEQ for Windows Version 3.0
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100.0%; Pred. No. 4.9e-89;
ive 0; Mismatches 0; Indels
75.4%; Score 365; DB 3; 1
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ive 0; Mismatches 0;
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365; Conservative
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ORGANISM: Homo sapien
                al Similarity
365; Conserv
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; GENERAL INFORMATION:
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RESULT 15
US-09-233-149A-115
Sequence 115, Application US/09232149A
Fatent No. 6465611
GENERAL INFORMATION:
APPLICANT: Witcham, Jangchun
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.427G
CURRENT APPLICATION NUMBER: US/09/232,149A
SOUTHAND SEQ ID NOS: 338
SOUTHAND SEQ ID NOS: 338
SEQ ID NO 115
LENGTH: 366
MANNER: PASISEQ FOR Windows Version 3.0
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ORGANISM: Homo sapien
US-09-232-149A-115
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Search completed: September 13, 2003, 03:05:40 Job time : 57 secs

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2. / cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
3. / cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
3. / cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
3. / cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
3. / cgn2_6/ptodata/2/pubpna/PCTUB_PUBCOMB.seq:*
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
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SUMMARIES

## Sequence 434, App Sequence 258, App Sequence 334, App Sequence 335, App Sequence 434, App Sequence 434, App Sequence 434 Sequence 434 Description US-09-895-793-434 US-09-895-814-434 US-09-895-814-434 US-10-012-896-434 US-10-010-940-434 US-10-025-300-258 US-10-025-300-258 US-09-759-143-335 US-09-780-669-335 US-09-822-827-335 US-09-822-827-335 US-09-822-827-335 US-09-835-880-335 US-09-832-880-335 US-09-895-814-335 US-09-759-143-434 US-09-780-669-434 US-09-822-827-434 100 100 100 100 100 100 Query Match Length DB 100.0 Score 484 484 484 484 484 484 473 472 473 U

335, App 335, App 335, App 282, App 102, App	103, App 223, App 222, App 115, App	115, A 115, A 115,	4 7 7 7 7 7 7	141, APP 141, APP 141, APP 141, APP 141, APP 141, APP 141, APP	
Sequence Sequence Sequence Sequence	000	Sequence Sequence Sequence Sequence Sequence	Sequence Sequence Sequence Sequence Sequence	Sequence 1 Sequence 1 Sequence 1 Sequence 2 Sequence 2 Sequence Sequence Sequence	
US-10-144-678A-335 US-10-012-896-335 US-10-010-940-335 US-10-205-823-282 US-10-202-193-102	US-10-202-193-103 US-10-202-193-223 US-10-202-193-222 US-09-759-143-115 US-09-780-669-115	US-09-02-03-115 US-09-115-453-115 US-09-232-880-115 US-09-895-793-115	S-09-89 S-10-14 S-10-01 S-10-01 S-09-96	US-09-780-669-141 US-09-780-669-141 US-09-822-827-141 US-09-115-453-141 US-09-895-793-141 US-09-895-793-141 US-09-895-880-141 US-09-895-880-141 US-10-144-678A-141	
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## ALIGNMENTS

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APPLICANT: Day, Craig H.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Carter, Darrick
APPLICANT: Li, Samuel
APPLICANT: Li, Samuel
APPLICANT: Skeiky, Vasir A.W.
APPLICANT: Wang, Aljun
APPLICANT: Mang, Aljun
APPLICANT: Belky, Vasir A.W.
APPLICANT: APPLICANT: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: COMPOSITIONS OF PROSTATE CANCER
FILE REPERBENCE: 210121.427C23
CURRENT APPLICATION NUMBER: US/09/759.143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
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100.0%; Pred. No. 7.7e-119;
tive 0; Mismatches 0;
               Sequence 434, Application US/09759143
Patent No. US2002002248A1
GENERAL INFORMATION:
                                                                                                                                           Harlocker, Susan L.
Jiang, Yuqui
Henderson, Robert A.
Kalos, Michael D.
                                                                               Xu, Jiangchun
Dillon, Davin C.
Mitcham, Jennifer L.
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Retter, Marc W.
Stolk, John A.
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Best Local Similarity 100.
Matches 484; Conservative
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US-09-759-143-434
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           CAGCCTGTTTCTATCCTGTTTAATAAATTAGTTTGGGTTCTTACATGCATAACAAACCC
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121,427C24
CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT FILING DATE: 2001-02-09
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SOFTWARE: FastSEQ for Windows Version 3.0
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Patent No. US20020051977A1
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Dillon, Davin C.
Mitcham, Jennifer L.
Harlocker, Susan L.
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Henderson, Robert A.
Kalos, Michael D.
Fanger, Gary R.
Retter, Marc W.
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McNeill, Patricia D.
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Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick
Li, Samuel
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Skeiky, Yasir A.W.
Hepler, William
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ORGANISM: Homo sapiens
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210.12.53411
CURRENT APPLICATION NUMBER: US/09/822,827
CURRENT APPLICATION NUMBER: 05/01.03-28
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 434
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    Score 484; DB 9; I
Pred. No. 7.7e-119;
0; Mismatches 0;
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Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 484; Conservative C
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Matches 484; Conserv
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; GENERAL INFORMATION:
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: COMPOSITIONS OF PROSTATE CANCER
FILLE REFERENCE: 210121.427C26
CURRENT APPLICATION NUMBER: US/09/895,814
CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 990
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 434
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Vinals de Bassols, Carlota
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Skeiky, Yasir A.W.
Hepler, William T.
Henderson, Robert A.
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Jiang, Yuqiu
Kalos, Michael D.
Retter, Marc W.
Stolk, John A.
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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Vedvick, Thomas S.
Carter, Darrick
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McNeill, Patricia
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Publication No.
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                  TITITCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAAAATCTGAAG
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APPLICANT: Mitcham, Journifer L.
APPLICANT: Mitcham, Journifer L.
APPLICANT: Kalos, Mitchal D.
APPLICANT: Retter, Marc W.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
APPLICANT: Wang, Aijun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Mitchal X.
APPLICANT: Monell, Dannel X.
APPLICANT: Henderson, Robert A.
APPLICANT: Houghton, Raymond L.
APPLICANT: Houghton, Raymond L.
APPLICANT: Houghton, Raymond L.
APPLICANT: Houghton, Raymond L.
APPLICANT: Foy, Teresa
APPLICANT: OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: COMPOSITIONS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C2
CURRENT APPLICATION NUMBER: US/09/895,793
CURRENT FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FESTESEQ for Windows Version 3.0
SEQ ID NO 434
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100.0%; Pred. No. 7.7e-119;
ive 0; Mismatches 0;
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APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Jiang, Yuqiu
APPLICANT: Kalos, Michael D.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Carter, Darrick
APPLICANT: Li, Samuel X.
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Matches 484; Conservative
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ORGANISM: Homo sapiens
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Pred. No. 7.7e-119;
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                     APPLICANT: Deng, Ta
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR T
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C28
CURRENT APPLICATION NUMBER: US/10/144,678A
CURRENT FILING DATE: 2002-08-12
NUMBER OF SEQ ID NOS: 1033
SOFTWARE: FASTERO for Windows Version 3.0
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Publication No. US20020183251A1
GENERAL INFORMATION:
APPLICANT: Vu, Jiangchun
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Jiang, Yuqiu
APPLICANT: Kalos, Michael D.
APPLICANT: Stolk, John A.
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APPLICANT: Vetter, Marc W.
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Matches 484; Conservative
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ORGANISM: Homo sapiens
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                                                                   tch 100.0%; Score 484; DB 10; al Similarity 100.0%; Pred. No. 7.7e-119; 484; Conservative 0; Mismatches 0;
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Houghton, Raymond L.
Vinals y de Bassols, Carlota
Foy, Teresa M.
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APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Hicham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Henderson, Robert A.
APPLICANT: Kalos, Michael D.
APPLICANT: Ranger, Gary R.
APPLICANT: Retter, Marc W.
APPLICANT: Retter, Marc W.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Garter, Jarck A.
APPLICANT: Li, Samuel X.
APPLICANT: Li, Samuel X.
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            ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-895-814-434
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Sequence 258, Application US/09925300 Patent No. US20020151681A1
                     Harlocker, Susan Louise
Jiang Yugui
Reed, Steven G.
Kalos, Michael
Mitcham, Jennifer L
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Matches 484; Conservative
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US-10-010-940-434
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       APPLICANT: Well AND APPLICANT: Skeiky, Yasir. APPLICANT: Skeiky, Yasir. APPLICANT: Hepler, William T. APPLICANT: Hepler, William T. APPLICANT: Henderson, Robert A. APPLICANT: Hural, John APPLICANT: Houghton, Raymond L. APPLICANT: Houghton, Raymond L. APPLICANT: Vinals de Bassols, Carlota APPLICANT: Foy, Teresa APPLICANT: Foy, Teresa APPLICANT: Wantanabe, Yoshihiro APPLICANT: Wantanaber, Yoshihiro APPLICANT: Wantanaber: 1016000515009 PROSTATE CANCER FILE REPERENCE: 210121.477.27 CURRENT APPLICATION NUMBER: US/10/012,896 CURRENT FILING DATE: 2001-12-10 NUMBER OF SEQ ID NOS: 1011 SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 434 LENDER OF SEQ ID NOS: 1014 LENDE
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100.0%; Pred. No. 7.7e-119;
tive 0; Mismatches 0;
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; GENERAL INFORMATION:
; APPLICANT: Xu, Jiangchun
; APPLICANT: Dillon, Davin C.
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Best Local Similarity 100.
Matches 484; Conservative
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; ORGANISM: Homo sapiens
US-10-012-896-434
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APPLICANT: Fanger, Gary
APPLICANT: Retter, Mark
APPLICANT: Solver, John
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REPERENCE: 210121.42703
CURRENT FILING DATE: 2001-12-05
SOFTWARE: FastSEQ for Windows Version 3.0
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APPLICANT: Steve Ruben
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA101
CURRENT APPLICATION NUMBER: US/09/925,300
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100.0%; Pred. No. 7.7e-119;
tive 0; Mismatches 0;
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                  FILE REFERENCE: PP-01532.103/200130.463D1
CURRENT APPLICATION NUMBER: US/10/202,193
CURRENT FILING DATE: 2002-07-23
NUMBER OF ESQ ID NOS: 341
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 334
LENGTH: 2051
                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (1)...(2051)
OTHER INFORMATION: n = A,T,C or G
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SENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
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Dillon, Davin C.
Mitcham, Jennifer L.
Harlocker, Susan L.
Jiang, Yuqui
Henderson, Robert A.
Ralos, Michael D.
Fanger, Gary R.
Fetter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
                                                                                                            TYPE: DNA
ORGANISM: Homo sapien
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|TCTTTA 369
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APPLICANT: Zhang, Jimmy
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll III, Eddie
APPLICANT: Endege, Wilson O.
APPLICANT: Monahan, John E.
APPLICANT: Schlegel, Robert
APPLICANT: Stellegel, Robert
APPLICANT: Stellegel, Robert
TITLE OF INVENTION: GENES AND GENE EXPRESSION PRODUCTS THAT
TITLE OF INVENTION: ARE DIFFERENTIALLY REGULATED IN PROSTATE CANCER
                                                                                                                                                                                                                                                                1;
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Best Local Similarity 99.8%; Pred. No. 8e-116;
Matches 484; Conservative 0; Mismatches
CURRENT FILING DATE: 2001-08-10
PRICR APPLICATION NUMBER: PCT/US00/05988
PRICR FILING DATE: 2000-03-08
PRICR FILING DATE: 1990-03-12
PRICR FILING DATE: 1990-03-12
NUMBER OF SEQ ID NOS: 1890
SEQ ID NO 258
                                                                                                                                                            ; NAME/KEY: misc feature
: LOCATION: (755)
; OTHER INDERARIION: n equals a,t,g, or
US-09-925-300-258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 334, Application US/10202193
Publication No. US20020192699A1
GENERAL INFORMATION:
                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
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2653 GCTTTTTCCCCCATTGGAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAACATCTGA 2712
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                                                                            ...rrlCant: Hural, John

APPLICANT: Hural, John

APPLICANT: McMeill, Patricia D.

APPLICANT: McWeill, Patricia D.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

FILE REPERDEC: 210121.427024

CURRENT APPLICATION NUMBER: US/09/780,669

CURRENT FILING DATE: 2001-02-09

NUMBER OF SEQ ID NOS: 943

SEPO-TET.
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Sequence 335, Application US/09022027

Sequence 335, Application US/09022027

SEQUENCE INFORMATION:

GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 2984;
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                          Day, Craig H.
Vedvick, Thomas
Carter, Darrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches 484; Conservative
          Stolk, John A.
                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: DNA
; ORGANISM: Homo sapien
US-09-780-669-335
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Best Local Similarity
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LENGTH: 2984
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APPLICANT: Carter, Darrick
APPLICANT: Li, Samuel
APPLICANT: Wang, Alling
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C23
CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF ESO ID NOS: 934
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                       Score 472; DB 9;
Pred. No. 2.9e-115;
0; Mismatches 0;
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Patent No. US20020051977A1
GENERAL INFORMATION:
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Dillon, Davin C.
Mitcham, Jennifer L.
Harlocker, Susan L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jiang, Yuqui
Henderson, Robert A.
Kalos, Michael D.
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Best Local Similarity 99.6%;
Matches 484; Conservative (
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Retter, Marc W.
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US-09-759-143-335
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LENGTH: 2984
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                                                                                  Gaps
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                              Length 2984;
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.5342.
CURRENT APPLICATION NUMBER: US/09/895,793
CURRENT FILING DATE: 2001-06-29
                                                    Indels
                                                    0
                         Score 472; DB 10;
Pred. No. 2.9e-115;
0; Mismatches 0;
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McMeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
FOy, Teresa
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b. US20020192763A1
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Sequence 335,
Publication No. Us.
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
"U,ICANT: Mitcham, Jennifer L.
"": Harlocker, Susan L.
"": "Ang, Yuqiu
                      97.58;
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Kalos, Michael D.
Retter, Marc W.
Stolk, John A.
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Vedvick, Thomas S.
Carter, Darrick
                 Ouery Match 97 59
Best Local Similarity 99.69
Matches 484; Conservative
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Hepler, William T.
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                                                                                                                                                  Length 2984;
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APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Mitchan, Jeanifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNODIAGNOSIS OF
TITLE OF INVENTION: PROSTATE
FILE REPERENCE: 210121.42866
CURRENT APPLICATION NUMBER: US/09/232,880
                                                                                                                                        97.5%; Score 472; DB 9; Length 296
99.6%; Pred. No. 2.9e-115;
Live 0; Mismatches 0; Indels
FILE REFERENCE: 210121.534C1
CURRENT APPLICATION NUMBER: US/09/822,827
CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SEQ ID NO 335
LENGTH: 2984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEC ID NOS: 338
SOFTWARE: FastSEQ for Windows Version 3.0
SOFTWARE: PastSEQ for Windows Version 3.0
LENGTH: 2984
TYPE: DNA
ORGANISM: Homo sapien
US-09-232-880-335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 335, Application US/09232880 Publication No. US20020182596A1
                                                                                                                                    Query Match
Best Local Similarity 99.67
Matches 484; Conservative
                                                                                     TYPE: DNA
ORGANISM: Homo sapien
US-09-822-827-335
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2953 TCTTTA 2958
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                                                                                                                                                                                                           121 TGTTGC--AAAAAAAAAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTT
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                                                                                  Length 2984;
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                                                                                 Score 472; DB 10;
Pred. No. 2.9e-115;
0; Mismatches 0;
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NUMBER OF SEQ ID NOS: 982
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 335
LENGTH: 2984
                                                                                 97.58;
                                                                                Query Match 97.5
Best Local Similarity 99.6
Matches 484; Conservative
                                       TYPE: DNA
CORGANISM: Homo sapien
US-09-895-793-335
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| TCTTTA 2958
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